



JC17 Rec'd PCT/PTO 21 MAY 2001

U.S. DEPARTMENT OF COMMERCE PATENT AND TRADEMARK OFFICE

ATTORNEY'S DOCKET NUMBER

UEMURA 5

**TRANSMITTAL LETTER TO THE UNITED STATES
DESIGNATED/ELECTED OFFICE (DO/EO/US)
CONCERNING A FILING UNDER 35 U.S.C. 371**

U.S. APPLICATION NO. (If known, see 37 CFR 1.5)

09/856319

INTERNATIONAL APPLICATION NO.
PCT/JP99/06473INTERNATIONAL FILING DATE
19 November 1999PRIORITY CLAIMED
20 November 1998

TITLE OF INVENTION

NOVEL SERINE PROTEASE BSSP5

APPLICANT(S) FOR DO/EO/US

Hidetoshi UEMURA et al.

Applicant herewith submits to the United States Designated/Elected Office (DO/EO/US) the following items and other information:

1. ☒ This is a **FIRST** submission of items concerning a filing under 35 U.S.C. 371.
2. ☐ This is a **SECOND** or **SUBSEQUENT** submission of items concerning a filing under 35 U.S.C. 371.
3. ☒ This is an express request to begin national examination procedures (35 U.S.C. 371(f)) at any time rather than delay examination until the expiration of the applicable time limit set in 35 U.S.C. 371(b) and PCT Articles 22 and 39(1).
4. ☒ The US has been elected in a Demand by the expiration of 19 months from the priority date (PCT Article 31).
5. ☒ A copy of the International Application as filed (35 U.S.C. 371(c)(2))
 - a. ☐ is attached hereto (required only if not transmitted by the International Bureau).
 - b. ☒ has been communicated by the International Bureau.
 - c. ☐ is not required, as the application was filed in the United States Receiving Office (RO/US).
6. ☒ An English language translation of the International Application as filed (35 U.S.C. 371(c)(2)).
7. ☒ Amendments to the claims of the International Application under PCT Article 19 (35 U.S.C. 371(c)(3))
 - a. ☐ are transmitted herewith (required only if not transmitted by the International Bureau).
 - b. ☐ have been communicated by the International Bureau.
 - c. ☐ have not been made; however, the time limit for making such amendments has NOT expired.
 - d. ☒ have not been made and will not be made.
8. ☐ An English language translation of the amendments to the claims under PCT Article 19 (35 U.S.C. 371(c)(3)).
9. ☒ An oath or declaration of the inventor(s) (35 U.S.C. 371(c)(4)).
10. ☐ An English language translation of the annexes to the International Preliminary Examination Report under PCT Article 36 (35 U.S.C. 371(c)(5)).

Items 11. to 16. below concern document(s) or information included:

11. ☐ An Information Disclosure Statement under 37 CFR 1.97 and 1.98.
12. ☐ An Assignment document for recording. A separate cover sheet in compliance with 37 CFR 3.28 and 3.31 is included.
13. ☒ A FIRST preliminary amendment.
 - ☐ A SECOND or SUBSEQUENT preliminary amendment.
14. ☐ A substitute specification.
15. ☐ A change of power of attorney and/or address letter.
16. ☒ Other items or information:

- ☒ Courtesy copy of the first page of the International Publication (WO 00/31243).
- ☒ Courtesy copy of the International Preliminary Examination Report. (In Japanese)
- ☒ Formal drawings, 6 sheets, Figures 1-6.
- ☒ Courtesy Copy of the International Search Report.
- ☒ Application Data Sheet

☒ The application is (or will be) assigned to: FUSO PHARMACEUTICAL INDUSTRIES, LTD. whose address is 7-10, Doshomachi 1-chome, Chuo-ku, Osaka-shi, Osaka Japan

U.S. APPLICATION NO. (If known, see 37 CFR 1.5)

097/856319

International Application No.

PCT/JP99/06473

Attorney's Docket No.

UEMURA 5

17. [xx] The following fees are submitted:

BASIC NATIONAL FEE (37 CFR 1.492 (a)(1)-(5):

Neither international preliminary examination fee (37 CFR 1.482)
nor international search fee (37 CFR 1.445(a)(2)) paid to USPTO
and International Search Report not prepared by the EPO or JPO.....**\$1000.00**

International preliminary examination fee (37 CFR 1.482) not paid to
USPTO but International Search Report prepared by the EPO or JPO.....**\$860.00**

International preliminary examination fee (37 CFR 1.482) not paid to USPTO but
international search fee (37 CFR 1.445(a)(2)) paid to USPTO**\$710.00**

International preliminary examination fee paid to USPTO (37 CFR 1.482)
but all claims did not satisfy provisions of PCT Article 33(1)-(4).....**\$690.00**

International preliminary examination fee paid to USPTO (37 CFR 1.482)
and all claims satisfied provisions of PCT Article 33(1)-(4).....**\$100.00**

CALCULATIONS PTO USE ONLY**ENTER APPROPRIATE BASIC FEE AMOUNT =**

\$ 860.00

Surcharge of **\$130.00** for furnishing the oath or declaration later than [] 20 [] 30
months from the earliest claimed priority date (37 CFR 1.492(e)).

\$

Claims as Originally Presented

Number Filed

Number Extra

Rate

Total Claims

81 - 20

61

X \$18.00

\$1,098.00

Independent Claims

12 - 3

09

X \$80.00

\$ 720.00

Multiple Dependent Claims (if applicable)

+\$270.00

\$ 270.00

TOTAL OF ABOVE CALCULATIONS =

\$2,948.00

Claims After Post Filing Prel. Amend

Number Filed

Number Extra

Rate

Total Claims

- 20

X \$18.00

\$

Independent Claims

- 3

X \$78.00

\$

TOTAL OF ABOVE CALCULATIONS =

\$2,948.00

Reduction of ½ for filing by small entity, if applicable. Applicant claims small entity
status. See 37 CFR 1.27.

\$

SUBTOTAL =

\$2,948.00

Processing fee of **\$130.00** for furnishing the English translation later than [] 20 [] 30
months from the earliest claimed priority date (37 CFR 1.492(f)).

\$

TOTAL NATIONAL FEE =

\$2,948.00

Fee for recording the enclosed assignment (37 CFR 1.21(h)). The assignment must be
accompanied by an appropriate cover sheet (37 CFR 3.28, 3.31). **\$40.00** per property +

\$

TOTAL FEES ENCLOSED =

\$2,948.00

Amount to be:

\$

refunded

charged

\$

a. [] A check in the amount of \$ _____ to cover the above fees is enclosed.

b. [X] Credit Card Payment Form (PTO-2038), authorizing payment in the amount of \$2,948.00, is attached.

c. [] Please charge my Deposit Account No. **02-4035** in the amount of \$ _____ to cover the above fees.

A duplicate copy of this sheet is enclosed.

d. [XX] The Commissioner is hereby authorized to charge any additional fees which may be required, or credit any overpayment
to Deposit Account No. **02-4035**. A duplicate copy of this sheet is enclosed.

NOTE: Where an appropriate time limit under 37 CFR 1.494 or 1.495 has not been met, a petition to revive (37 CFR 1.137(a) or
(b)) must be filed and granted to restore the application to pending status.

SEND ALL CORRESPONDENCE TO:

BROWDY AND NEIMARK, P.L.L.C.
624 NINTH STREET, N.W., SUITE 300
WASHINGTON, D.C. 20001

TEL: (202) 628-5197

FAX: (202) 737-3528

Date of this submission: **May 21, 2001**

Form PTO-1350 (as slightly revised by Browdy and Neimark)

SIGNATURE

Roger L. Browdy

NAME

25,618

REGISTRATION NUMBER

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:)	Art Unit:
Hidetoshi UEMURA et al.)	
)	
IA No.: PCT/JP99/06473)	
)	Washington, D.C.
IA Filed: 19 November 1999)	
)	
U.S. App. No.:)	
(Not Yet Assigned))	May 21, 2001
)	
National Filing Date:)	
(Not Yet Received))	
)	
For: NOVEL SERINE PROTEASE BSSP5)		Docket No.: UEMURA 5

PRELIMINARY AMENDMENT

Honorable Commissioner for Patents
Washington, D.C. 20231

Sir:

Contemporaneous with the filing of this case and
prior to calculation of the filing fee, kindly amend as
follows:

IN THE SPECIFICATION

After the title please insert the following
paragraph:

REFERENCE TO RELATED APPLICATIONS

--The present application is the national stage
under 35 U.S.C. §371 of international application
PCT/JP99/06473, filed 19 November 1999 which designated the
United States, and which application was not published in the
English language.--

0956319-052101

In re of: Hidetoshi UEMURA et al. (UEMURA 5)

REMARKS

The above amendment to the specification is being made to insert reference to the PCT application of which the present case is a U.S. national stage.

Favorable consideration is earnestly solicited.

Respectfully submitted,
BROWDY AND NEIMARK, P.L.L.C.
Attorneys for Applicant

By: 

Roger L. Browdy
Registration No. 25,618

RLB:wrđ

Telephone No.: (202) 628-5197

Facsimile No.: (202) 737-3528

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101250 61295860



Rec'd PCT/PTO 15 AUG 2001 #3

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:)	Box PCT
UEMURA et al)	Examiner:
Appln. No.: 09/856,319)	Washington, D.C.
IA No.: PCT/JP99/06473)	
IA Filed: November 19, 1999)	August 15, 2001
For: NOVEL SERINE PROTEASE BSSP5))	Atty.Docket: UEMURA=5

RESPONSE TO NOTIFICATION TO COMPLY WITH
SEQUENCE LISTING REQUIREMENTS

Honorable Commissioner for Patents
Washington, D.C. 20231

Sir:

In response to the Notification to Comply dated June 22, 2001, and prior to the examination of the above-described application, please amend the present application as follows:

IN THE SPECIFICATION

Please replace the paragraph beginning at page 22, line 24 and ending at page 24, line 4, with the following rewritten paragraph:

--The vector is not specifically limited in so far as it can express the protein of the present invention. Examples thereof include pBAD/His, pRSETA, pcDNA2.1, pTrcHis2A, pYES2, pBlueBac4.5, pcDNA3.1 and pSecTag2 manufacture by Invitrogen, pET and pBAC manufactured by Novagen, pGEM manufactured by Promega, pBluescriptII manufactured by Stratagene, pGEX and pUC18/19 manufactured by

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Pharmacia, PfastBAC1 manufactured by GIBCO and the like. Preferably, a protein expression vector (described in the specification of a patent application entitled "Protein expression vector and its use" and filed by the same applicant on the same day) is used. This expression vector is constructed by using pCRII-TOPO vector described in the Examples hereinafter, or a commercially available expression vector, for example pSecTag2A vector or pSecTag2B vector (Invitrogen) and integrating a secretory signal nucleotide sequence suitable for expression of the protein of the present invention, in the 3' downstream side thereof, a Tag nucleotide sequence, a cleavable nucleotide sequence and a cloning site, into which a nucleotide sequence encoding a target protein can be inserted, in this order. More specifically, it is preferred to use trypsin signal as the secretory signal, a nucleotide sequence encoding polyhistidine as the Tag nucleotide sequence, and a nucleotide sequence encoding an amino acid sequence which is susceptible to enzyme-specific cleavage, i.e., a nucleotide sequence encoding the amino acid sequence of Asp-Asp-Asp-Asp-Lys (SEQ ID NO:33) (said amino acid sequence is recognized by enterokinase, and the recombinant fusion protein is cleaved at the C-terminus part thereof) as the cleavable nucleotide sequence.--

IN THE SEQUENCE LISTING

Please substitute the attached Sequence Listing, numbered as pages 1-13 for the one previously submitted.

REMARKS

Applicants have substituted into the present specification a new paper copy Sequence Listing section according to 37 C.F.R. §1.821(c) as new pages 1-13. Furthermore, attached hereto is a 3 1/2" disk containing the "Sequence Listing" in computer readable form in accordance with 37 C.F.R. §1.821(e).

Applicants have made changes to the general information section and have added to the previously filed Sequence Listing new SEQ ID NO:33 listed on page 24, line 1, and have amended the specification to insert SEQ ID NO:33, as supported in the present specification.

The following statement is provided to meet the requirements of 37 C.F.R. §1.825(a) and 1.825(b).

I hereby state, in accordance with 37 C.F.R. §1.825(a), that the amendments included in the substitute sheets of the sequence listing are believed to be supported in the application as filed and that the substitute sheets of the sequence listing are not believed to include new matter.

I hereby further state, in accordance with 37 C.F.R. §1.825(b), that the attached copy of the computer readable form is the same as the attached substitute paper copy of the sequence listing.

Under U.S. rules, each sequence must be classified in <213> as an "Artificial Sequence", a sequence of "Unknown" origin, or a sequence originating in a particular organism, identified by its scientific name.

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Neither the rules nor the MPEP clarify the nature of the relationship which must exist between a listed sequence and an organism for that organism to be identified as the origin of the sequence under <213>.

Hence, counsel may choose to identify a listed sequence as associated with a particular organism even though that sequence does not occur in nature by itself in that organism (it may be, e.g., an epitopic fragment of a naturally occurring protein, or a cDNA of a naturally occurring mRNA, or even a substitution mutant of a naturally occurring sequence). Hence, the identification of an organism in <213> should not be construed as an admission that the sequence per se occurs in nature in said organism.

Similarly, designation of a sequence as "artificial" should not be construed as a representation that the sequence has no association with any organism. For example, a primer or probe may be designated as "artificial" even though it is necessarily complementary to some target sequence, which may occur in nature. Or an "artificial" sequence may be a substitution mutant of a natural sequence, or a chimera of two or more natural sequences, or a cDNA (i.e., intron-free sequence) corresponding to an intron-containing gene, or otherwise a fragment of a natural sequence.

The Examiner should be able to judge the relationship of the enumerated sequences to natural sequences by giving full consideration to the specification, the art cited therein, any further art cited in an IDS, and the

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results of his or her sequence search against a database containing known natural sequences.

Attached hereto is a marked-up version of the changes made to the specification by the current amendment. The attached page is captioned "Version with markings to show changes made".

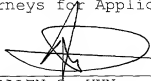
Applicants submit that the present application contains patentable subject matter and therefore urge the examiner to pass the case to issuance.

If the examiner has any questions or comments concerning the above described application, the examiner is urged to contact the undersigned at the phone number below.

Respectfully submitted,

BROWDY AND NEIMARK, P.L.L.C.
Attorneys for Applicant(s)

By



ALLEN C. YUN
Registration No. 37,971

ACY:al
624 Ninth Street, N.W.
Washington, D.C. 20001
Telephone No.: (202) 628-5197
Facsimile No.: (202) 737-3528

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VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the specification:

The paragraph beginning at line 24 of page 22 and ending at line 4 of page 24 has been amended as follows:

The vector is not specifically limited in so far as it can express the protein of the present invention. Examples thereof include pBAD/His, pRSETA, pcDNA2.1, pTrcHis2A, pYES2, pBlueBac4.5, pcDNA3.1 and pSecTag2 manufacture by Invitrogen, pET and pBAC manufactured by Novagen, pGEM manufactured by Promega, pBluescriptII manufactured by Stratagene, pGEX and pUC18/19 manufactured by Pharmacia, PfastBAC1 manufactured by GIBCO and the like. Preferably, a protein expression vector (described in the specification of a patent application entitled "Protein expression vector and its use" and filed by the same applicant on the same day) is used. This expression vector is constructed by using pCRII-TOPO vector described in the Examples hereinafter, or a commercially available expression vector, for example pSecTag2A vector or pSecTag2B vector (Invitrogen) and integrating a secretory signal nucleotide sequence suitable for expression of the protein of the present invention, in the 3' downstream side thereof, a Tag nucleotide sequence, a cleavable nucleotide sequence and a cloning site, into which a nucleotide sequence encoding a target protein can be inserted, in this order. More specifically, it is preferred to use trypsin signal as the secretory signal, a nucleotide sequence encoding polyhistidine as the Tag nucleotide sequence, and a nucleotide sequence encoding an amino acid sequence which is susceptible to

enzyme-specific cleavage, i.e., a nucleotide sequence encoding the amino acid sequence of Asp-Asp-Asp-Asp-Lys (SEQ ID NO:33) (said amino acid sequence is recognized by enterokinase, and the recombinant fusion protein is cleaved at the C-terminus part thereof) as the cleavable nucleotide sequence.

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101250 612550

SEQUENCE LISTING

<110> UEMURA, Hidetoshi
 OKUI, Akira
 KOMINAMI, Katsuya
 YAMAGUCHI, Nozomi
 MITSUI, Shinichi

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035539 052101

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115 120 125

Ser Pro Ala Arg Tyr Thr Ala Gln Val Ser Pro Val Cys Leu Ala Ser
130 135 140

Thr Asn Glu Ala Leu Pro Ser Gly Leu Thr Cys Val Thr Thr Gly Trp
145 150 155 160

Gly Arg Ile Ser Gly Val Gly Asn Val Thr Pro Ala Arg Leu Gln Gln
165 170 175

Val Val Leu Pro Leu Val Thr Val Asn Gln Cys Arg Gln Tyr Trp Gly
180 185 190

Ala Arg Ile Thr Asp Ala Met Ile Cys Ala Gly Gly Ser Gly Ala Ser
195 200 205

Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Gln Lys Gly Asn
210 215 220

Thr Trp Val Leu Ile Gly Ile Val Ser Trp Gly Thr Lys Asn Cys Asn
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245 250 255

Ile Asn Gln Val Met Ala Tyr Asn
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00856210.052107

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 <210> 8
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 <210> 9
 <211> 35
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<213> Artificial Sequence

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<223> Designed oligonucleotide primer to amplify a portion of plasmid
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<210> 10

<211> 29

<212> DNA

<213> Artificial Sequence

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<223> Designed oligonucleotide primer to amplify a portion of plasmid
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<210> 11

<211> 26

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<210> 12

<211> 17

<212> DNA

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<210> 13

<211> 17

<212> DNA

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17

<210> 14
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serin proteases-encoding sequence

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20

<210> 16
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 <210> 17
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 <223> Designed oligonucleotide primer for RACE for hBSSP5 (forward)
 <400> 17
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 <210> 18
 <211> 20
 <212> DNA
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 <210> 19
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<210> 22
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<210> 23
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<400> 25
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 for mBSSP5 (reverse)

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Asp Asp Asp Asp Lys
 1 5

6/PRTS

NOVEL SERINE PROTEASE BSSP5

5

FIELD OF THE INVENTION

The present invention relates to isolated polynucleotides of human and mouse serine proteases (hereinafter referred to as "hBSSP5" and "mBSSP5", respectively, and, in case no differentiation thereof from each other is needed, simply referred to as "BSSP5"), and their homologous forms, mature forms, precursors and polymorphic variants as well as a method for detecting thereof. Further, it relates to hBSSP5 and mBSSP5 proteins, compositions containing hBSSP5 and mBSSP5 polynucleotides and proteins, as well as their production and use.

BACKGROUND OF THE INVENTION

In general, proteases are biosynthesized as inactive precursors. They undergo limited hydrolysis in molecules to convert into activated type proteases. In so far as enzymes are proteases, they have an activity for hydrolyzing a peptide bond, while their action modes are varied according to kinds of proteases. According to a particular kind of catalytic site, proteases are divided into serine proteases, cysteine proteases, aspartate

proteases, metal proteases and the like. Proteases of each kind have a variety of properties, ranging from a protease having general digestive properties to a protease having various regulatory domains and strict substrate specificity, thereby specifically hydrolyzing only characteristic proteins.

Further, proteins undergo various processing even after translation to produce active proteins. In many secretory proteins, a protein are first synthesized on the ribosome in cytoplasm as an inactive precursor (pro-form) which comprises an active protein bearing at the N-terminus thereof a peptide of about 15 to 60 amino acids responsible for secretion (secretory signal). This peptide region is concerned with the mechanism for passing through the cell membrane and is removed upon cleavage by a specific protease during the passage through the membrane, in almost all the cases, to produce the mature type protein. A secretory signal has a broad hydrophobic region comprising hydrophobic amino acids in the middle of the sequence, and basic amino acid residues at a site close to the N-terminus. A secretory signal is a synonym of a signal peptide. In addition, in some proteins, a peptide moiety which functions as a secretory signal is further attached to the N-terminus of the inactive precursor. Such a protein is called a prepro-protein (prepro-form).

For example, trypsin is present as a prepro-form immediately after translation into amino acids. After being secreted from cells, it is present as a pro-form and is converted into active trypsin in duodenum upon limited hydrolysis by enteropeptidase or by trypsin itself.

The optimal pH range of serine proteases is neutral to weak alkaline and, in general, many of them have a molecular weight of about 30,000 or lower. All proteases of blood coagulation, fibrinolysis and complement systems having a large molecular weight belong to trypsin-like serine proteases. They have many regulator domains and form a protease cascade which is of very importance to reactions in a living body.

Recently, cDNAs and amino acid sequences of many novel proteases have been determined by PCR for consensus sequences of serine proteases using oligonucleotide primers. According to this method, novel proteases have been found by various researchers such as Yamamura et al. (Yamanura, Y et al., Biochem. Biophys. Res. Commun., 239, 386, 1997), Gschwend, et al. (Gschwend, T. P. et al., Mol. Cell. Neurosci., 9, 207, 1997), Chen et al. (Chen, Z-L, et al., J. Neurosci., 15, 5088, 1995) and others.

SEQ ID NO: 3 of JP 9-149790 A discloses neurosin as a novel serine protease. Neurosin has also been reported in Biochimica et Biophysica Acta, 1350, 11-14,

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1997. By this, there is provided a method for mass production of neurosin using the serine protease gene and a method for screening specific inhibitors using the enzyme. In addition, the screening method has been shown to be
5 useful for screening medicines for treating various diseases.

Serine proteases expressed in a brain-nerve system such as neurosin are considered to play various roles in the brain-nerve system. Therefore, there is a
10 possibility that isolation of a gene encoding a novel protease expressed in a brain-nerve system and production of a protein using the gene would be useful for diagnosis or treatment of various diseases related to the brain-nerve system.

15 Nowadays, in general, clinical diagnosis of Alzheimer's disease is conducted based on the diagnosis standard of DSM-III-R and NINCDS-ADRDA (Mckhann, G. et al., Neurology, 34. 939, 1994) or the diagnosis standard of DSM-IV (American Psychiatric Association; Diagnostic and
20 statistical manuals of mental disorders, 4th ed., Washington DC, American Psychiatric Association, 1994). However, these standards are conditioned by decline of recognition functions which causes a severe disability in a daily life or a social life. Then, it is pointed out that
25 the diagnosis is less scientific objectivity because the

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diagnosis may be influenced by the level of an individual's social life and further the specialty and experience of a physician who diagnoses particular conditions. In addition, definite diagnosis of Alzheimer's disease is conducted by pathohistological analyses and, in this respect, substantial inconsistency between clinical diagnosis and autopsy diagnosis is pointed out.

At present, image diagnosis is employed as a supplemental means in clinical diagnosis of Alzheimer's diagnosis and it is possible to analyze brain functions, for example, decline of metabolism and atrophy in specific sites such as hippocampus, parietal lobe of cerebral cortex and the like which are specific for Alzheimer's disease by PET and SPECT. However, to define Alzheimer's disease based on lowering of a blood flow from parietal lobe to temporal lobe is very dangerous. In addition, there is few report showing that MRS testicle useful for patients with dementia including those of Alzheimer's disease. Further, although CT-MRI image diagnosis is used, a lesion of white matter such as atrophy of brain, PVL or the like is not specific for Alzheimer type dementia. Since it has been reported that atrophy of brain proceeds as getting older, the above observation is not necessarily found in Alzheimer type dementia. Furthermore, since an image obtained by MRI varies according to strength of a magnetic field,

performance of an apparatus and imaging conditions, numerical data obtain in different facilities cannot be compared with each other except atrophic change. In addition, there is a limit to image measurement. Further, enlargement of ventricle can be recognized in vascular dementia cases and there are cases wherein atrophy of hippocampus is observed after ischemia of basilar artery.

Under these circumstances, many researchers have requested to develop biological diagnosis markers as a means for providing better precision and objectivity for clinical diagnosis of Alzheimer's disease. At the same time, the following important roles in the future will be expected.

1) Objective judgment system of effect of medicaments for treating Alzheimer's disease.

2) Detection of Alzheimer's disease before a diagnosis standard is met, or disease conditions are manifested.

Further, data obtained in different facilities can be compared with each other by using the same diagnosis marker. Therefore, development of biological diagnosis markers is recognized to be a most important field among fields of Alzheimer's disease studies and its future prospects will be expected. Approaches to development of biological diagnosis markers up to now are divided into

that based on constitute components of characteristic pathological changes of Alzheimer's disease such as senile plaque and neurofibril change, and an approach based on other measures. Examples of the former include

5 cerebrospinal fluid tau protein, A β and its precursor, β APP. Examples of the latter include mydriasis test with cholinergic drug, Apo E and other genes relating to Alzheimer's disease. However, no good results are obtained.

Serine proteases are also considered to play

10 important role in cancer cells. The reason why extermination of cancer by surgical treatment or topical irradiation of radioactive ray is difficult is metastasis capability of cancer. For spread of solid tumor cells in a body, they should loosen their adhesion to original

15 adjacent cells, followed by separating from an original tissue, passing through other tissues to reach blood vessel or lymph node, entering into the circulatory system through stratum basal and endothelial layer of the vessel, leave from the circulatory system at somewhere in the body, and

20 surviving and proliferating in a new environment. While adhesion to adjacent epidermal cells is lost when expression of cadherin which is an intercellular adhesive molecule of epithelium is stopped, to break through tissues is considered to depend on proteolytic enzymes which

25 decompose an extracellular matrix.

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As enzymes which decompose the matrix, mainly, metal proteases (Rha, S. Y. et al., Breast Cancer Research Treatment, 43, 175, 1997) and serine proteases are known. They cooperate to decompose matrix protein such as collagen, laminin and fibronectin. Among serine proteases known to be concerned in decomposition of the matrix, in particular, there is urokinase type plasminogen activator (U-PA). U-PA has a role as a trigger specific for a protein decomposition chain reaction. Its direct target is plasminogen. It is present in blood abundantly and is a precursor of an inactive serine protease which accumulates in reconstructed sites of tissues such as injured sites and tumors as well as inflammatory sites. In addition, as proteases which are concerned in metastasis and infiltration of cancers, for example, a tissue factor, lysosomal type hydrolase and collagenase have been known.

At present, cancer is the top cause of death in Japan and more than 200,000 people are died per year. Then, specific substances which can be used as markers for diagnosis and therapy or prophylaxis of cancer are studied intensively. Such specific substances are referred to as tumor markers or tumor marker relating biomarkers. They are utilized in aid of diagnosis before treatment of cancer, for presuming carcinogenic organ and pathological tissue type, for monitoring effect of treatment, for finding

recurrence early, for presuming prognosis, and the like. At present, tumor markers are essential in clinical analyses. Among them, alpha fetoprotein (AFP) which has high specificity to hepatocellular carcinoma and yolk sac tumor (Taketa K. et al., Tumour Biol., 9, 110, 1988), and carcinoembryonic antigen (CEA) are used worldwide. In the future, tumor markers will be required more and more, and it is desired to develop, for example, organ specific markers and tumor cell specific markers which are highly reliable serologic diagnosis of cancer. Up to now, humunglandular kallikrein (hK2) which is a serine protease expressed at human prostatic epithelial cells has been reported as a marker for prostatic cancer. And, hK2 has 78% homology with the sequence of prostatic specific antigen (PSA) and PSA is also used widely as a biochemical marker of prostatic cancer (Mikolajczyk, S. d. et al., Prostate, 34, 44, 1998; Pannek, J. et al., Oncology, 11, 1273, 1997; Chu, T. M. et al., Tumour Biology, 18, 123, 1997; Hsieh, M. et al., Cancer Res., 57, 2651, 1997).

20

OBJECTS OF THE INVENTION

Thus, the main object of the present invention is to provide a novel serine protease which can be used for treating or diagnosing various diseases such as Alzheimer's disease (AD), epilepsy, cancer, inflammation, sterility,

25

prostate hypertrophy and the like in various tissues such as brain, lung, prostate, testicle, skeletal muscle, liver and the like, and can be used as an excellent marker instead of that presently used.

5

SUMMARY OF THE INVENTION

Under these circumstances, the present inventors have succeeded in cloning of cDNA encoding novel human and mouse serine proteases.

10

In summary, the 1st feature of the present invention is amino acid sequences of biological active mature serine proteases hBSSP5 and mBSSP5 and nucleotide sequences encoding the amino acid sequences.

15

That is, they are the amino acid sequence composed of 231 amino acids (mature type hBSSP5 (the 1st to 231th amino acids of SEQ ID NO: 2)) and a nucleotide sequence encoding the amino acid sequence (the 110th to 802nd bases of SEQ ID NO: 1). In addition, they include amino acid sequences substantially similar to SEQ ID NO: 2 and nucleotide sequences encoding such similar amino acid sequences. Further, they include modified derivatives of proteins having these amino acid sequences. An amino acid sequence substantially similar to a given amino acid sequence used herein means an amino acid sequence derived from the given amino acid sequence by modification such as

25

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substitution, deletion, addition and/or insertion of one to several amino acids with maintaining the same property as that of the protein having the given amino acid sequence. The modified derivative of the proteins includes, for example, phosphate adduct, sugar chain adduct, metal adduct (e.g., calcium adduct), the protein fused to another protein such as albumin etc., dimer of the protein, and the like.

Further, they are the amino acid sequence composed of 231 amino acids (mature type mBSSP5 (the 1st to 231th amino acids of SEQ ID NO: 4)) and a nucleotide sequence encoding the amino acid sequence (the 132nd to 824th bases of SEQ ID NO: 3). In addition, they include amino acid sequences substantially similar to the anubi acid sequence and nucleotide sequences encoding such similar amino acid sequences. Further, they include modified derivatives of proteins having these amino acid sequences.

The 2nd feature of the present invention is an amino acid sequence composed of 33 amino acids represented by the -33rd to -1st amino acids represented by SEQ ID NO: 4 and a nucleotide sequence encoding the amino acid sequence (the 33rd to 131st bases of SEQ ID NO: 3). In addition, this feature includes amino acid sequences substantially similar to the amino acid sequence and

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nucleotide sequences encoding these substantially similar amino acid sequences. Further, this feature includes modified derivatives of proteins having these amino acid sequences.

5 The 3rd feature of the present invention is an amino acid sequence composed of 264 amino acids (precursor type mBSSP5 (the -33rd to 231st amino acids of SEQ ID NO: 4)) wherein 33 amino acids compsed of -33rd to -1st amino acids represented by SEQ ID NO: 4 are added to the N-terminus side of the mature type mBSSP5 amino acid sequence (SEQ ID NO: 4) and a nucleotide sequence encoding the amino acid sequence (the 33rd to 824th bases of SEQ ID NO: 3). In addition, this feature includes amino acid sequences substantially similar to the amino acid sequence represented by SEQ ID NO: 4 and nucleotide sequences encoding these substantially similar amino acid sequences. Further, this feature includes modified derivatives of proteins having there amino acid sequences.

10

15

The present invention also relates to the nucleotide sequences represented by SEQ ID NOS: 1 and 3 and nucleotide sequences similar to these sequences.

20

The 4th feature of the present invention is a vector comprising the nucleotide sequence according to any of the above 1st to the 3rd features, and transformant cells transformed with the vector.

25

The 5th feature of the present invention is a process for producing BSSP5 protein from the transformed cells of the 4th feature.

5 The 6th feature of the present invention is a transgenic non-human animal, wherein the expression level of BSSP5 gene has been altered.

The 7th feature of the present invention is an antibody against BSSP5 protein or its fragment and a process for producing thereof.

10 The 8th feature of the present invention is a method for determining BSSP5 protein or its fragment in a specimen using the antibody of the 7th feature.

The 9th feature of the present invention is a diagnostic marker of diseases comprising BSSP5 protein.

15 The 10th feature of the present invention is a method for detecting pancreatitis by determining concentration of BSSP5 protein, a pharmaceutical composition comprising an antibody against BSSP5 protein or its fragment, and use of BSSP5 protein for preparing an
20 antibody for detecting pancreatitis.

Hereinafter, unless otherwise stated, the nucleotide sequence represented by each SEQ ID NO: includes the above-described various fragments thereof, and similar nucleotide sequences and their fragments. Likewise, the
25 amino acid sequence represented by each SEQ ID NO: includes

conventional manner. For preparing mRNA, guanidine isothiocyanate-calcium chloride method (Chirwin, et al., Biochemistry, 18, 5294, 1979) or the like can be used. For preparing poly (A) + RNA from total RNAs, there can be used

5 affinity chromatography using a carrier, for example, Sepharose, latex particles, etc., to which oligo (dT) is attached, and the like. The above-obtained RNA can be used as a template and treated with reverse transcriptase by using, as a primer, oligo (dT) which is complementary to

10 the poly (A) strand at the 3'-terminus, or a random primer, or a synthesized oligonucleotide corresponding to a part of the amino acid sequence of hBSSP5 or mBSSP5 to obtain a hybrid mRNA strand comprising DNA complementary to the mRNA or cDNA. The double stranded DNA can be obtained by

15 treating the above-obtained hybrid mRNA strand with *E. coli* RNase, *E. coli* DNA polymerase and *E. coli* DNA ligase to convert into a DNA strand.

It is also possible to carry out cloning by RT-PCR method using primers synthesized on the basis of the

20 nucleotide sequence of hBSSP5 or mBSSP5 gene and using hBSSP5 or mBSSP5 expressing cell poly (A) + RNA as a template. Alternatively, the desired cDNA can be obtained without using PCR by preparing or synthesizing a probe on the basis of the nucleotide sequence of hBSSP5 or mBSSP5

25 gene and screening a cDNA library directly. Among genes

obtained by these methods, the gene of the present invention can be selected by confirming a nucleotide sequence thereof. The gene of the present invention can also be prepared according to a conventional method using chemical syntheses of nucleic acids, for example, phosphoamidite method (Mattenccci, M. D. et al., J. Am. Chem. Soc., 130, 3185, 1981) and the like.

By using the thus-obtained hBSSP5 or mBSSP5 gene, their expression in various tissues can be examined.

In case of northern blotting analysis, the expression of hBSSP5 is observed in pancreas and the expression of mBSSP5 is observed in spleen. In case of RT-PCR analysis, the expression of hBSSP5 is observed in brain of the fetuses and placenta of the adults, and mBSSP5 shows the expression in brain of newborn to grown-up mice and in testicle of grown-up mice. Then, the novel proteases of the present invention are presumed to play various roles in brain, placenta, testicle, pancreas and spleen. For example, in brain, there is a possibility that they can be used for treatment and diagnosis of brain diseases such as Alzheimer's disease (AD), epilepsy, brain tumor and the like. Further, in other tissues, there is a possibility that BSSP5 of the present invention and a gene encoding it can be used for treatment and diagnosis of various diseases such as cancer, inflammation, sterility, prostate

hypertrophy and the like. Further, it is presumed they may have a certain influence on blood coagulation, fibrinolysis and complement systems. Furthermore, there is a possibility that inhibitors of serine proteases can be used for treatment and prevention of Alzheimer's disease, epilepsy, cancer, inflammation, sterility, prostate hypertrophy and the like. Moreover, increase in a blood level of BSSP5 of the present invention is observed in a rat pancreatitis model and therefore it can be used for detection of pancreatitis.

The present inventors have shown that the mature type of novel human serine protease (hBSSP5) is composed of 231 amino acids, and the mature type of novel mouse serine protease (mBSSP5) is composed of 231 amino acids and its precursor type is composed of 264 amino acids. Further, the amino acid sequences of the mature type serine proteases contain consensus sequences having serine protease activity.

The term "pro part" used herein means a part of a pro-form, i.e., the pro-form from which the corresponding active type protein part is removed. The term "pre part" used herein means a part of a prepro-form, i.e., the prepro-form from which the corresponding pro-form is removed. The term "prepro part" used herein means a part of a prepro-form, i.e., the prepro-form from which the

corresponding active type protein part is removed.

The amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2 is the BSSP5 mature or active type protein composed of 231 amino acids, and the nucleotide sequence encoding the amino acid sequence represented by the 110th to 802nd bases of SEQ ID NO: 1 is composed of 693 bases. The present inventors have shown that the serine protease activity is maintained even when one to several amino acids of the N-terminus in the amino acid sequence of the mature type protein of hBSSP5 is deleted or added, while the sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2 is preferred.

The amino acid sequence represented by SEQ ID NO: 4 is mBSSP5 protein composed of 264 amino acids, and the nucleotide sequence encoding the amino acid sequence represented SEQ ID NO: 3 is composed of 792 bases. The present inventors have shown that the serine protease activity is maintained even when one to several amino acids of the N-terminus in the amino acid sequence of the mature type protein of mBSSP5 is deleted or added, while the sequence represented by SEQ ID NO: 4 is preferred. The sequence of the -33th to -1st amino acids of SEQ ID NO: 4 is the prepro or pro part and the amino acid sequence represented by the -33rd to 231th of the amino acids is considered to be a precursor type of mBSSP5 protein.

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In general, many genes of eucaryote exhibit polymorphism and, sometimes, one or more amino acids are substituted by this phenomenon. Further, even in such case, sometimes, a protein maintains its activity. Then, the present invention includes a gene encoding a protein obtained by modifying a gene encoding the amino acid sequence represented by SEQ ID NO: 2 or 4, artificially, in so far as the protein has the characteristic function of the gene of the present invention. Further, the present invention includes a protein which is a modification of the amino acid sequence represented by SEQ ID NO: 2 or 4 in so far as the protein has the characteristics of the present invention. Modification is understood to include substitution, deletion, addition and/or insertion. In particular, the present inventors have shown that, even when several amino acids are added to or deleted from the N-terminus amino acid of hBSSP5 or mBSSP5 mature protein represented by SEQ ID NO: 2 or 4, the resultant sequence maintains its activity.

That is, the present invention includes a protein comprising either amino acid sequence described in SEQ ID NOS: 2 and 4; or one of these amino acid sequences wherein one to several amino acids have been substituted, deleted, added and/or inserted, and being belonging to serine protease family.

Each codon for the desired amino acid itself has been known and it can be selected freely. For example, codons can be determined according to a conventional manner by taking into consideration of frequency of use of codons in a host to be utilized (Grantham, R. et al., Nucleic Acids Res., 9, 43, 1989). Therefore, the present invention also includes a nucleotide sequence appropriately modified by taking into consideration of degeneracy of a codon. Further, these nucleotide sequences can be modified by a site directed mutagenesis using a primer composed of a synthetic oligonucleotide encoding the desired modification (Mark, D. F. et al., Proc. Natl. Acad. Sci. USA., 81, 5662, 1984), or the like.

Furthermore, the DNA of the present invention includes DNA which is hybridizable to either of nucleotide sequences described in SEQ ID NOS: 1 and 3, or nucleotide sequences complementary to these nucleotide sequences in so far as the protein encoded by the nucleotide sequence has the same properties as those of hBSP5 or mBSP5 of the present invention. It is considered that many of sequences which are hybridizable to a given sequence under stringent conditions have a similar activity to that of a protein encoded by the given sequence. The stringent conditions according to the present invention includes, for example, incubation in a solution containing 5 x SSC, 5% Denhardt's

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solution (0.1% BSA, 0.1% Ficoll 1400, 0.1% PVP), 0.5% SDS
and 20 µg/ml denatured salmon sperm DNA at 37°C overnight,
followed by washing with 2 x SSC containing 0.1% SDS at
room temperature. Instead of SSC, SSPE can be
5 appropriately used.

Probes for detecting a hBSSP5 or mBSSP5 gene can
be designed based on either of nucleotide sequences
described in SEQ ID NOS: 1 and 3. Or, primers can be
designed for amplifying DNA or RNA containing the
10 nucleotide sequence. To design probes or primers is
carried out routinely by a person skilled in the art. An
oligonucleotide having a designed nucleotide sequence can
be synthesized chemically. And, when a suitable label is
added to the oligonucleotide, the resultant oligonucleotide
15 can be utilized in various hybridization assays. Or, it
can be utilized in nucleic acid synthesis reactions such as
PCR. An oligonucleotide to be utilized as a primer has,
preferably, at least 10 bases, more preferably 15 to 50
bases in length. An oligonucleotide to be utilized as a
20 probe has, preferably, 100 bases to full length.

Moreover, it is possible to obtain a promoter
region and an enhancer region of a hBSSP5 or mBSSP5 gene
present in the genome based on the cDNA nucleotide sequence
of hBSSP5 or mBSSP5 provided by the present invention.
25 Specifically, these control regions can be obtained

Examples thereof include pBAD/His, pRSETA, pcDNA2.1, pTrcHis2A, pYES2, pBlueBac4.5, pcDNA3.1 and pSecTag2 manufacture by Invitrogen, pET and pBAC manufactured by Novagen, pGEM manufactured by Promega, pBluescriptII

5 manufactured by Stratagene, pGEX and pUC18/19 manufactured by Pharmacia, PfastBAC1 manufactured by GIBCO and the like. Preferably, a protein expression vector (described in the specification of a patent application entitled "Protein

10 expression vector and its use" and filed by the same applicant on the same day) is used. This expression vector is constructed by using pCRII-TOPO vector described in the Examples hereinafter, or a commercially available expression vector, for example pSecTag2A vector or pSecTag2B vector (Invitrogen) and integrating a secretory

15 signal nucleotide sequence suitable for expression of the protein of the present invention, in the 3' downstream side thereof, a Tag nucleotide sequence, a cleavable nucleotide sequence and a cloning site, into which a nucleotide sequence encoding a target protein can be inserted, in this

20 order. More specifically, it is preferred to use trypsin signal as the secretory signal, a nucleotide sequence encoding polyhistidine as the Tag nucleotide sequence, and a nucleotide sequence encoding an amino acid sequence which is susceptible to enzyme-specific cleavage, i.e., a

25 nucleotide sequence encoding the amino acid sequence of

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Asp-Asp-Asp-Asp-Lys (said amino acid sequence is recognized by enterokinase, and the recombinant fusion protein is cleaved at the C-terminus part thereof) as the cleavable nucleotide sequence.

5 Furthermore, the present invention provides transformed cells having the nucleotide sequence of the present invention in an expressible state by means of the above vector. Preferably, host cells to be used for the transformed cells of the present invention are animal cells and insect cells. However, host cells include any cells (including those of microorganisms) which can express a nucleotide sequence encoding the desired protein in the expression vector of the present invention and can secrete extracellularly.

15 The animal cells and insect cells used herein
include cells derived from human being and cells derived
from fly or silk worm. For example, there are CHO cell,
COS cell, BHK cell, Vero cell, myeloma cell, HEK293 cell,
HeLa cell, Jurkat cell, mouse L cell, mouse C127 cell,
20 mouse FM3A cell, mouse fibroblast, osteoblast, cartilage
cell, S2, Sf9, Sf21, High Five™ (registered trade mark)
cell and the like. The microorganisms used herein include
E. coli, yeast or the like.

The protein of the present invention as such can
25 be expressed as a recombinant fused protein so as to

facilitate isolation, purification and recognition. The recombinant fused protein used herein means a protein expressed as an adduct wherein a suitable peptide chain are added to the N-terminus and/or C-terminus of the desired protein expressed by a nucleotide sequence encoding the desired protein. The recombinant protein used herein means that obtained by integrating a nucleotide sequence encoding the desired protein in the expression vector of the present invention and cut off an amino acid sequence which derived from nucleic acids other than those encoding the desired protein from the expressed recombinant fused protein, and is substantially the same as the protein of the present invention.

Introduction of the above vector into host cells can be carried out by, for example, transfection according to lipopolyamine method, DEAE-dextran method, Hanahan method, lipofectin method or calcium phosphate method, microinjection, eletroporation and the like.

As described above, the present invention also relates to a process for producing hBSSP5 or mBSSP5 comprising culturing cells transformed with the above nucleotide sequence of the present invention and collecting the produced hBSSP5 or mBSSP5. The culture of cells and separation and purification of the protein can be carried out by a per se known method.

The present invention also relates to an inhibitor of the novel serine protease of the present invention. Screening of the inhibitor can be carried out according to a per se known method such as comparing the enzyme activity upon bringing into contact with a candidate compound with that without contact with the candidate compound, or the like

The present invention relates to a non-human transgenic animal whose expression level of hBSSP5 or mBSSP5 gene has been altered. The hBSSP5 or mBSSP5 gene used herein includes cDNA, genomic DNA or synthetic DNA encoding hBSSP5 or mBSSP5. In addition, expression of a gene includes any steps of transcription and translation. The non-human transgenic animal of the present invention is useful for studies of functions or expression control of hBSSP5 or mBSSP5, elucidation of mechanisms of diseases in which hBSSP5 or mBSSP5 is presumed to be involved, and development of disease model animals for screening and safety test of medicine.

In the present invention, expression of a gene can be modified artificially by mutagenizing at a part of several important sites which control normal gene expression (enhancer, promoter, intron, etc.) such as deletion, substitution, addition and/or insertion to increase or decrease an expression level of the gene in

comparison with its inherent expression level. This mutagenesis can be carried out according to a known method to obtain the transgenic animal.

In a narrow sense, the transgenic animal means an animal wherein a foreign gene is artificially introduced into reproductive cells by gene recombinant techniques. In a broad sense, the transgenic animal includes an antisense transgenic animal the function of whose specific gene is inhibited by using antisense RNA, an animal whose specific gene is knocked out by using embryonic stem cells (ES cells), and an animal into which point mutation DNA is introduced, and the transgenic animal means an animal into which a foreign gene is stably introduced into a chromosome at an initial stage of ontogeny and the genetic character can be transmitted to the progeny.

The transgenic animal used herein should be understood in a broad sense and includes any vertebrates other than a human being. The transgenic animal of the present invention is useful for studies of functions or expression control of hBSSP5 or mBSSP5, elucidation of mechanisms of diseases associated with cells expressing in a human being, and development of disease model animals for screening and safety test of medicine.

As a technique for creating the transgenic animal, a gene is introduced into a nucleus in a pronucleus stage

of egg cells with a micropipette directly under a phase-contrast microscope (microinjection, U.S. Patent 4,873,191). Further, there are a method using embryonic stem cell (ES cell), and the like. In addition, there are newly
5 developed methods such as a method wherein a gene is introduced into a retroviral vector or adenoviral vector to infect egg cells, a sperm vector method wherein a gene is introduced into egg cells through sperms, and the like.

A sperm vector method is a gene recombinant
10 technique wherein a foreign gene is incorporated into sperm cells by adhesion, electroporation, etc., followed by fertilization of egg cells to introduce the foreign gene into the egg cells (M. Lavitrano et al., Cell, 57, 717, 1989). Alternatively, an in vivo site specific gene
15 recombinant technique such as that using cre/loxP recombinase system of bacteriophage P1, FLP recombinase system of *Saccharomyces cerevisiae*, etc. can be used. Furthermore, introduction of a transgene of the desired protein into a non-human animal using a retroviral vector
20 has been reported.

For example, a method for creating a transgenic animal by microinjection can be carried out as follows.

First, a transgene primarily composed of a promoter responsible for expression control, a gene
25 encoding a specific protein and a poly A signal is required.

It is necessary to confirm expression modes and amounts between respective systems because an expression mode and amount of a specific molecule is influenced by a promoter activity, and transgenic animals differ from each other according to a particular system due to the difference in a copy number of an introduced transgene and a introduction site on a chromosome. An intron sequence which is spliced may be previously introduced before the poly A signal because it has been found that an expression amount varies due to a non-translation region and splicing. Purity of a gene to be used for introduction into fertilized egg cells should be as high as possible. This is of importance. Animals to be used include mice for collecting fertilized eggs (5- to 6-week-old), male mice for mating, false pregnancy female mice, seminiferous tubule-ligated mice, and the like.

For obtaining fertilized egg cells efficiently, ovulation may be induced with gonadotropin or the like. Fertilized egg cells are recovered and a gene in an injection pipette is injected into male pronucleus of the egg cells by microinjection. For returning the injected egg cells to a fallopian tube, an animal (false pregnancy female mouse, etc.) is provided and about 10 to 15 eggs/mouse are transplanted. Then, genomic DNA is extracted from the end part of the tail to confirm whether

wherein all of either of the allele genes have been modified or destroyed. Further, a homozygote mouse can be created by mating heterozygote mice.

Homologous recombination means recombination
5 between two genes whose nucleotide sequences are the same or very similar to each other in terms of gene recombination mechanism. PCR can be employed to select homologous recombinant cells. A PCR reaction can be carried out by using a part of a gene to be inserted and a
10 part of a region where the insertion is expected as primers to find out occurrence of homologous recombination in cells which give an amplification product. Further, for causing homologous recombination in a gene expressed in embryonic stem cells, homologous recombinant cells can readily be
15 selected by using a known method or its modification. For example, cells can be selected by joining a neomycin resistant gene to a gene to be introduced to impart neomycin resistance to cells after introduction.

The present invention also provide an antibody
20 recognizing hBSSP5 or mBSSP5 or a fragment thereof. The antibody of the present invention includes an antibody against a protein having the amino acid sequence described in SEQ ID NO: 2 or 4 or its fragment. An antibody against hBSSP5 or mBSSP5 or a fragment thereof (e.g., polyclonal
25 antibody, monoclonal antibody, peptide antibody) or an

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antiserum can be produced by using hBSSP5 or mBSSP5 or a fragment thereof, etc. as an antigen according to a per se known process for producing an antibody or an antiserum.

5 The hBSSP5 or mBSSP5 or a fragment thereof is administered to a site of a warm-blooded animal where an antibody can be produced by administration thereof as such or together with a diluent or carrier. For enhancing the antibody production, upon administration, Freund's complete adjuvant or Freund's incomplete adjuvant may be
10 administrated. Normally, the administration is carried out once every 1 to 6 weeks, 2 to 10 times in all. Examples of the warm-blooded to be used include monkey, rabbit, dog, guinea pig, mouse, rat, sheep, goat, chicken and the like with mouse and rat being preferred. As rats, for example,
15 Wistar and SD rats are preferred. As mice, for example, BALB/c, C57BL/6 and ICR mice are preferred.

For producing monoclonal antibody producer cells, individuals whose antibody titer have been recognized are selected from warm-blooded animals, e.g., a mouse immunized
20 with an antigen. Two to 5 days after the last immunization, the spleen or lymph node of the immunized animal is collected and antibody producer cells contained therein are subjected to cell fusion with myeloma cells to prepare a monoclonal antibody producer hybridoma. The antibody titer
25 in an antiserum can be determined by, for example, reacting

the antiserum with a labeled hBSSP5 or mBSSP5 as described hereinafter, followed by measurement of the activity bound to the antibody. The cell fusion can be carried out according to a known method, for example, that described by Koehler and Milstein (Nature, 256, 495, 1975) or its modifications (J. Immunol. Method, 39, 285, 1980; Eur. J. biochem, 118, 437, 1981; Nature, 285, 446, 1980). As a fusion promoting agent, there are polyethylene glycol (PEG), Sendai virus and the like. Preferably, PEG is used. Further, for improving fusion efficiency, lectin, poly-L-lysine or DMSO can be appropriately added.

Examples of myeloma cells include X-63Ag8, NS-1, P3U1, SP2/0, AP-1 and the like with SP2/0 being preferred. The preferred ratio of the number of the antibody producer cells (spleen cells) : the number of myeloma cells are 1 : 20 to 20 : 1. PEG (preferably PEG 1000 to PEG 6000) is added at a concentration of about 10 to 80% and the mixture is incubated at 20 to 40°C, preferably 30 to 37°C for 1 to 10 minutes to carry out the cell fusion efficiently. Screening of anti-hBSSP5 or mBSSP5 antibody producer hybridomas can be carried out by various methods. For example, a supernatant of a hybridoma culture is added to a solid phase to which hBSSP5 or mBSSP5 antigen is adsorbed directly or together with a carrier (e.g., microplate), followed by addition of an anti-immunoglobulin antibody (in

case that the cells used in cell fusion is those of a mouse, anti-mouse immunoglobulin antibody is used) or protein A to detect the anti-hBSSP5 or mBSSP5 monoclonal antibody attached to the solid phase. Or, a supernatant of a hybridoma culture is added to a solid phase to which an anti-immunoglobulin antibody or protein A is adsorbed, followed by addition of hBSSP5 or mBSSP5 labeled with a radioactive substance, an enzyme, etc., to detect the anti-hBSSP5 or mBSSP5 monoclonal antibody attached to the solid phase.

Selection and cloning of the anti-hBSSP or mBSSP monoclonal antibody can be carried out according to a per se known method or its modification. Normally, a HAT (hypoxanthine, aminopterin, thymidine)-added medium for culturing animal cells is used. Any culture medium can be used for selection, cloning and growing up in so far as the hybridoma can grow. For example, there can be used RPMI culture medium containing 1 to 20%, preferably 10 to 20% fetal bovine serum, or a serum-free medium for culturing hybridomas. Preferably, the culture is carried out at a temperature of about 37°C. Normally, the culture time is 5 days to 3 weeks, preferably 1 weeks to 2 weeks. Normally, the culture is carried out under 5% CO₂. The antibody titer of a supernatant of a hybridoma culture can be measured according to the same manner as that of the above-

described measurement of anti-BSSP5 antibody titer in an antiserum. That is, examples of the measurement to be used include radioimmunoassay (RIA), enzyme-linked immunosorbent assay (ELISA), FIA (fluorescence immunoassay), plaque assay, 5 agglutination reaction method, and the like. Among them, ELISA as shown blew is preferred.

Screening by ELISA

A protein prepared according to the same operation as that for an immunogen is immobilized on the 10 surface of each well of an ELISA plate. Next, BSA, MSA, OVA, KLH, gelatin, skimmed milk, or the like is immobilized on each well to prevent non-specific adsorption. A supernatant of a hybridoma culture is added to each well and is allowed to stand for a given time so that an 15 immunological reaction proceeds. Each well is washed with a washing solution such as PBS or the like. Preferably, a surfactant is added to this washing solution. An enzyme labeled secondary antibody is added and allowed to stand for a given time. As the enzyme to be used for the label, 20 there can be used β -galactosidase, alkaline phosphatase, peroxidase and the like. After washing each well with the same washing solution, a substrate solution of the labeled enzyme used is added so that an enzymatic reaction proceeds. When the desired antibody is present in the supernatant of 25 a hybridoma culture, the enzymatic reaction proceeds and

the color of the substrate solution is changed.

Normally, cloning is carried out by a per se known method such as semi-solid agar method, limiting dilution method and the like. Specifically, after
5 confirming a well in which the desired antibody is produced by the above-described method, cloning is carried out to obtain a single clone. For cloning, it is preferred to employ limiting dilution method wherein hybridoma cells are diluted so that one colony is formed per one well of a
10 culture plate. For cloning by limiting dilution method, feeder cells can be used, or a cell growth factor such as interleukin 6, etc. can be added to improve colony forming capability. In addition, cloning can be carried out by using FACS and single cell manipulation method. The cloned
15 hybridoma is preferably cultured in a serum-free culture medium and an optimal amount of an antibody is added to its supernatant. The single hybridoma thus obtained can be cultured in a large amount by using a flask or a cell culture device, or cultured in the abdominal cavity of an
20 animal (J. Immunol. Meth., 53, 313, 1982) to obtain a monoclonal antibody. When culturing in a flask, there can be used a cell culture medium (e.g., IMDM, DMEM, RPMI1640, etc.) containing 0 to 20% of FCS. When culturing in the abdominal cavity of an animal, the animal to be used is
25 preferably the same species or the same line as that from

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which the myeloma cells used in the cell fusion are derived, a thymus deficient nude mouse or the like, and the hybridoma is transplanted after administration of a mineral oil such as pristane, etc. After 1 to 2 weeks, myeloma
5 cells are proliferated in the abdominal cavity to obtain ascites containing a monoclonal antibody.

The monoclonal antibody of the present invention which does not cross-react with other proteins can be obtained by selecting a monoclonal antibody which
10 recognizes an epitope specific to hBSSP5 or mBSSP5. In general, an epitope presented by an amino acid sequence composed of at least 3, preferably 7 to 20 successive amino acid residues in an amino acid sequence which constitutes a particular protein is said to be an inherent epitope of the
15 protein. Then, a monoclonal antibody recognizing an epitope constituted by a peptide having an amino acid sequence composed of at least 3 successive amino acid residue selected from the amino acid residues disclosed in either of SEQ ID NOS: 2 and 4 can be said to be the
20 monoclonal antibody specific for hBSSP5 or mBSSP5 of the present invention. An epitope common to BSSP5 family can be selected by selecting an amino acid sequence conservative among the amino acid sequences described in SEQ ID NOS: 2 and 4. Or, in case of a region containing an
25 amino acid sequence specific for each sequence, a

monoclonal antibody which can differentiate respective proteins can be selected.

Separation and purification of the anti-hBSSP5 or mBSSP5 monoclonal antibody, like a conventional polyclonal antibody, can be carried out according to the same manner as those of immunoglobulins. As a known purification method, there can be used a technique, for example, salting out, alcohol precipitation, isoelectric precipitation, electrophoresis, ammonium sulfate precipitation, absorption and desorption with an ion exchange material (e.g., DEAE), ultrafiltration, gel filtration, or specific purification by collecting only an antibody with an antibody-binding solid phase or an active adsorber such as protein A or protein G, etc., and dissociating the binding to obtain the antibody. For preventing formation of aggregates during purification or decrease in the antibody titer, for example, human serum albumin is added at a concentration of 0.05 to 2%. Alternatively, amino acids such as glycine, α -alanine, etc., in particular, basic amino acids such as lysine, arginine, histidine, etc., saccharides such as glucose, mannitol, etc., or salts such as sodium chloride, etc. can be added. In case of IgM antibody, since it is very liable to be aggregated, it may be treated with β -propiolactone and acetic anhydride.

The polyclonal antibody of the present invention

can be produced according to a per se known method or its modification. For example, an immunogen (protein antigen) per se or a complex thereof with a carrier protein is prepared and, according to the same manner as that in the

5 above monoclonal antibody production, a warm-blooded animal is immunized. A material containing an antibody against the protein of the present invention or its fragment is collected from the immunized animal and the antibody is separated and purified to obtain the desired antibody. As

10 for a complex of an immunogen and a carrier protein for immunizing a warm-blooded animal, the kind of a carrier protein and the mixing ratio of a carrier and a hapten are not specifically limited in so far as an antibody against the hapten immunized by cross-linking with the carrier is

15 efficiently produced. For example, there can be used about 0.1 to 20, preferably about 1 to 5 parts by weight of bovine serum albumin, bovine cycloglobulin, hemocyanin, etc. coupled with one part by weight of a hapten. For coupling a carrier and a hapten, various condensing agents can be

20 used. Examples thereof include glutaraldehyde, carbodiimide or maleimide active ester, active ester agents having thiol group or dithiopyridyl group, and the like. The condensed product is administered as such or together with a carrier or diluent to a site of a warm-blooded

25 animal where an antibody can be produced. For enhancing

the antibody production, upon administration, Freund's complete adjuvant or Freund's incomplete adjuvant may be administrated. Normally, the administration is carried out once every 2 to 6 weeks, 3 to 10 times in all. The polyclonal antibody can be collected from blood, ascites, or the like, preferably blood of the immunized animal. The polyclonal antibody titer in an antiserum can be measured according to the same manner as measurement of the above monoclonal antibody titer in the antiserum. Separation and purification of the polyclonal antibody, like the above monoclonal antibody, can be carried out according to the same manner as those of immunoglobulins.

The monoclonal antibody and polyclonal antibody against hBSSP5 or mBSSP5 or a fragment thereof can be utilized for diagnosis and treatment of diseases associated with cells expressing hBSSP5 or mBSSP5. By using these antibodies, hBSSP5 or mBSSP5 or a fragment thereof can be determined based on their immunological binding to hBSSP5 or mBSSP5 or a fragment thereof of the present invention. Specifically, examples of a method for determining hBSSP5 or mBSSP5 or a fragment thereof by using these antibodies include a sandwich method wherein the antibody attached to an insoluble carrier and the labeled antibody are reacted with hBSSP5 or mBSSP5 or a fragment thereof to form a sandwich complex and the sandwich complex is detected, as

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well as a competitive method wherein labeled hBSSP5 or mBSSP5, and hBSSP5 or mBSSP5 or a fragment thereof in the specimen are competitively reacted with the antibody and hBSSP5 or mBSSP5 or a fragment thereof in the specimen is determined based on the amount of the labeled antigen reacted with the antibody.

As a sandwich method for determining hBSSP5 or mBSSP5 or a fragment thereof, there can be used two step method, one step method and the like. In two step method, first, the immobilized antibody is reacted with hBSSP5 or mBSSP5 or a fragment thereof and then unreacted materials are completely removed by washing, followed by addition of the labeled antibody to form immobilized antibody-hBSSP5 or mBSSP5-labeled antibody. In one step method, the immobilized antibody, labeled antibody and hBSSP5 or mBSSP5 or a fragment thereof are added at the same time.

Examples of an insoluble carrier used for the determination include synthetic resins such as polystyrene, polyethylene, polypropylene, polyvinyl chloride, polyester, polyacrylate, nylon, polyacetal, fluorine plastic, etc.; polysaccharides such as cellulose, agarose, etc.; glass; metal; and the like. An insoluble carrier may be shaped in various forms, for example, tray, sphere, fiber, rod plate, container, cell, test tube, and the like. The antibody adsorbed by a carrier is stored at a cold place in the

presence of an appropriate preservative such as sodium azide or the like.

For immobilization of the antibody, a known chemical bonding method or a physical adsorption can be used. Examples of a chemical bonding method include a method using glutaraldehyde; maleimide method using N-succusinimidyl-4-(N-maleimidomethyl)cyclohexane-1-carboxylate, N-succusinimidyl-2-maleimide acetate or the like; carbodiimide method using 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide hydrochloride; or the like. In addition, there are maleimidobenzoyl-N-hydroxysuccinimide ester method, N-succinimidyl-3-(2-pyridylthio)propionic acid method, bisdiazobenzidine method, and dipalmityllysine method. Or, it is possible to capture a complex formed beforehand by reacting a material to be tested with two antibodies, whose epitopes are different, with an immobilized a 3rd antibody against the antibody.

For labeling, it is preferred to use enzyme, fluorescent substance, luminous substance, radioactive substance, metal chelate, or the like. Examples of the enzyme include peroxidase, alkaline phosphatase, β -D-galactosidase, malate dehydrogenase, *Staphylococcus* nuclease, δ -5-steroidisomerase, α -glycerol phosphate dehydrogenase, triose phosphate isomerase, horseradish peroxidase, asparaginase, glucose oxidase, ribonuclease,

urease, catalase, glucose-6-phosphate dehydrogenase, glucoamylase, acetylcholinesterase and the like. Examples of the fluorescent substance include fluorescein isothiocyanate, phycobiliprotein, rhodamine, phycoerythrin, 5 phycocyanin, allophycocyanin, o-phthalaldehyde, and the like. Examples of the luminous substance include isoluminol, lucigenin, luminol, aromatic acridinium ester, imidazole, acridinium salt and its modified ester, luciferin, luciferase, aequorin and the like. Examples of the 10 radioactive substance include ^{125}I , ^{127}I , ^{131}I , ^{14}C , ^3H , ^{32}P , ^{35}S and the like. The labeling material is not limited to them and any material which can be used for immunological determination can be used. Further, a low molecular weight hapten such as biotin, dinitrophenyl, pyridoxal or 15 fluorescamine may be attached to the antibody. Preferably, horseradish peroxidase is used as a labeling enzyme. This enzyme can be reacted with various substrates and can readily be attached to the antibody by periodate method.

When an enzyme is used as a labeling material, a 20 substrate and, if necessary, a coloring enzyme is used for measuring its activity. In case of using peroxidase as the enzyme, H_2O_2 is used as a substrate and, as a coloring agent, there can be used 2,2'-azino-di-[3-ethylbenzthiazoline sulfonic acid] ammonium salt (ABTS), 25 5'-aminosalicylic acid, o-phenylenediamine, 4-

aminoantipyrine, 3,3',5,5'-tetramethylbenzidine and the like. In case of using alkaline phosphatase as the enzyme, o-nitorphenylphosphate, p-nitrophenylphosphoric acid, or the like can be used as a substrate. In case of using β -D-galactosidase as the enzyme, fluorescein-d-(β -D-galactopyranoside), 4-methylumbelliphenyl- β -D-galactopyranoside, or the like can be used as a substrate. The present invention also include a kit comprising the above monoclonal antibody, polyclonal antibody and reagents.

As a cross-linking agent, a known cross-linking agent such as N,N'-o-phenylenedimaleimide, 4-(N-maleimidomethyl)cyclohexanoate N-succinimide ester, 6-maleimidohexanoate N-succineimide ester, 4,4'-dithiopyridine or the like can be utilized. The reaction of these cross-linking agents with enzymes and antibodies can be carried out by a known method according to properties of a particular cross-linking agent. Further, as the antibody, a fragment thereof, for example, Fab', Fab, F(b'2) can be used as the case may be. A labeled enzyme can be obtained by the same treatment regardless of whether the antibody is polyclonal or monoclonal. When the above labeled enzyme obtained by using a cross-linking agent is purified by a known method such as affinity chromatography or the like, a immunoassay system having more higher sensitivity can be obtained. The enzyme labeled and

as primers. Namely, 5 μ l of the template, 5 μ l of 10 x ExTaq buffer, 5 μ l of dNTP, 10 pmol of each of the above primers and 0.5 μ l of ExTaq (TAKARA) were added and the total volume was adjusted to 50 μ l with sterilized water.

5 PCR was carried out by repeating a cycle of heating at 94°C for 0.5 minute, at 55°C for 0.5 minute and then at 72°C for 1 minutes, 35 times. The PCR product was mixed with pCR II-TOPO vector attached to TOPO TA cloning kit (Invitrogen) and the mixture was allowed to stand at room temperature

10 for 5 minutes. Then, according to a conventional manner, *E. coli* Top 10 attached to the kit was transformed and applied to a LB (Amp^r) plate (containing 100 μ g/ml of ampicillin). According to a conventional manner, a plasmid was extracted from each colony obtained and its nucleotide sequence was

15 determined by cycle sequencing method with a fluorescence sequencer (ABI). Homology of the sequence of each clone was examined by means of GenBank. Regarding an unknown sequence, i.e., BSSP5 gene, the full length cDNA was obtained by 5' RACE and 3' RACE and, according to the same

20 manner as described above, the nucleotide sequence was determined. Namely, BSSP5 clone specific primers, GSP1 primers (primers having nucleotide sequences of SEQ ID NOS: 16 and 18) and GSP2 primers [primers having SEQ ID NOS: 17 and 19) were prepared. PCR was carried out by using human

25 brain Marathon-Ready cDNA (Clontech), AP1 primer attached

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to this reagent and the above GSP1 primer and heating at 94°C for 2 minutes once and repeating a cycle of heating at 94°C for 30 seconds, at 60°C for 30 seconds and then at 72°C for 30 seconds 35 times. Then, 5 µl of the PCR product diluted to 1/100, 5 µl of 10 x buffer, 5 µl of dNTP, 10 pmol of either of 10 µM of the above GSP2 primer, 10 pmol of AP2 primer attached to the above reagent and 0.5 unit of ExTaq were admixed and adjusted to 50 µl with sterilized water. Then, according to the same manner as the above, PCR was carried out. The PCR product was cloned by the above TOPO TA cloning kit and sequenced to obtain the upstream and downstream regions of the above clone. Further, based on this sequence, the primers capable of amplifying ORF [hBSSP5F1 (SEQ ID NO: 20), hBSSP5R1/E (SEQ ID NO: 22)] were prepared and PCR carried out using human brain Marathon-ready cDNA as a template to confirm that these clones were identical. This was cloned into pCR II-TOPO vector attached to TOPO TA cloning kit to obtain the plasmid pCR II/hBSSP5 containing the full length cDNA clone. The nucleotide sequence of DNA contained in this plasmid is shown in SEQ ID NO: 1 and the amino acid sequence of hBSSP5 protein deduced from the nucleotide sequence is shown in SEQ ID NO: 2.

According to the same manner, the plasmid pCRII/mBSSP5 containing a mouse homologous gene was

obtained by carrying out 5' RACE and 3' RACE using mouse brain Marathon-Ready cDNA (Clontech) as a template, followed by cloning. The nucleotide sequence contained in this plasmid is shown in SEQ ID NO: 3 and the amino acid sequence deduced from this nucleotide sequence is shown in SEQ ID NO: 4. The amino acid sequence shown in SEQ ID NO: 4 is mBSSP5 protein composed of 264 amino acids and the nucleotide sequence encoding this protein shown in SEQ ID NO 3 is composed of 792 bases. The sequence represented by the -33rd to -1st amino acids of SEQ ID4: is a prepro or pro part and the amino acid sequence represented by the -33rd to 231st amino acids is considered to be a precursor of mBSSP5 protein.

	SEQ ID	Name of primer	Direction	Sequence	Use
15	15	15	15	15	15
					NO:
					human BSSP5
	16		Forward	TGTCAGCCCTGGCCGCCATT	RACE
	17		Forward	GCGAGTATGACCGATCATCA	RACE
20	18		Reverse	CGCCACCTGCACAGATCATG	RACE
	19		Reverse	GAATCAGTGCCGGCAGTACT	RACE
	20	hBSSP5F1	Forward	TGCCACGATGTTGCTGCTCA	FL*
	21	hBSSP5F2	Forward	ATTGTCAACGGGGAGAATGC	mature
	22	hBSSP5R1/E	Reverse	GGAATTCGGGTCTTTAATGGGTTGAGC	FL*
25	23	hBSSP5R4	Reverse	CCTGGCACGAGGAGGCAC	for RT-PCR

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mousse BSSP5

	24	mBSSP5F1	Forward	ACCATGAACAATGACCTGAC	RACE
	25	mBSSP5F2	Forward	GAATCAGTGTCCGGCAGT	RACE
	26	mBSSP5F3	Forward	GACCATCTCAACACCATTCC	FL*
5	27	mBSSP5Fmature	Forward	ATTGTCAACGGGGAGAATGC	mature
	28	mBSSP5.1	Reverse	ATGGCATCGGTAATGCGTGC	RACE
	29	mBSSP5R2	Reverse	CAGGTGTTTCCCTTCTGGCA	RACE
	30	mBSSP5R3/E	Reverse	GGAATTCGGACAGTTTAGTTGTAGGCC	FL*

*: for full length

10 Example 2

Expression of hBSSP5 or mBSSP5 gene in human beings or mice internal organs

According to the protocol of QuickPrep Micro mRNA purification Kit (Amersham-Pharmacia), mRNAs were isolated from various internal organs of Balb/c mice or their fetuses. They were subjected to electrophoresis according to a conventional manner and transcribed to a nylon membrane. A probe was prepared separately by isolating a part of a nucleotide sequence encoding the mature protein of mBSSP5 (the 132nd to 824th bases of SEQ ID NO: 3) from pCR II/mBSSP5, purifying it and labeling it with α -³²P dCTP. The probe was diluted with 5 x SSC and reacted with the above membrane filter at 65°C for a whole day and night. According to the same manner, a probe was prepared by isolating a part of a nucleotide sequence encoding the

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mature protein of hBSSP5 (the 110th to 802nd bases of SEQ ID NO: 1) from pCR II/hBSSP5, purifying it and labeling it with α -³²P dCTP. The probe was diluted with 5 x SSC and reacted with human multiple tissue blot (Clontech) membrane at 65°C for a whole day and night. Then, each membrane filter was washed twice each with 2 x SSC/0.1% SDS at room temperature for 30 minutes, 1 x SSC/0.1% SDS at room temperature for 30 minutes and 0.1 x SSC/0.1% SDS at 65°C for 30 minutes. The filter was exposed to an imaging plate for FLA2000 (Fuji Film) for one day to analyze the expression. The results shown in the drawings are those obtained by using human multiple tissue blot (clontech) membrane (Fig. 1) and mRNAs prepared from various internal organs of 3-month-old mice (Fig. 2). In addition, the mRNAs prepared above were subjected to RT-PCR by using Ready To Go RT-PCR Beads (Amersham-Pharmacia) and hBSSP5 or mBSSP5 gene specific primers according to the protocol attached to the kit (amplification by using SEQ ID NOS: 20 and 22 and further amplification by using SEQ ID NOS: 21 and 23).

As seen from Figs. 1 and 2, in case of northern blotting analysis, the expression of hBSSP5 was observed in pancreas and the expression of mBSSP5 was observed in spleen. Further, in case of RT-PCR, the expression of hBSSP5 was observed in brain of fetuses and in placenta in

adults. The expression of mBSSP5 was observed in brain and testicle of fetuses to grown up mice. Then, it is presumed that the novel serine proteases have various roles in brain, placenta, pancreas, spleen and tenticle.

5 Example 3

Determination of enzyme activity of novel serine protease mature protein encoded by hBSSP5 or mBSSP5 gene

(1) Construction of expression plasmid

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10 A cDNA fragment containing the region encoding the mature protein of hBSSP5 or mBSSP5 protein was amplified by PCR using the plasmid pCR II/hBSSP5 or pCR II/mBSSP5 as a template (the primers used were SEQ ID NOS: 21 and 22 for human being, and SEQ ID NOS: 27 and 30 for mouse). Each PCR product was ligated to pTrc-HisB
15 (Invitrogen) which had been digested with BamHI and blunted with mung bean nuclease according to a conventional method. *E. coli* JM109 was transformed by the resultant and colonies formed were analyzed by PCR to obtain *E. coli* containing the desired serine protease expressing plasmid
20 pTrcHis/hBSSP5 or pTrcHis/mBSSP5.

The resultant *E. coli* strains were designated *E. coli* pTrcHis/hBSSP5 and *E. coli* pTrcHis/mBSSP5 and deposited at National Institute of Bioscience and Human-Technology (NIBH), Agency of Industrial Science &
25 Technology of 1-1-3 Higashi, Tsukuba-shi, Ibaraki-ken,

Japan on October 29, 1998 under the accession numbers of FERM P-17038 and FERM P-17035, respectively.

(2) Expression of protein by *E. coli* containing expression plasmid

5 A single colony of *E. coli* having the expression
plasmid was inoculated in 10 ml of LB (Amp^r) culture medium
and incubated at 37°C overnight. This was inoculated in
250 ml of LB (Amp^r) culture medium and incubated at 37°C.
When the absorbance at 600 nm became 0.5, 250 µl of 0.1 M
10 IPTG (isopropyl-β-D-(-)-thiogalactopyranoside) was added
and the incubation was continued for additional 5 hours.
The *E. coli* was centrifuged and suspended in a cell
disruption buffer (10 mM phosphate buffer pH 7.5, 1 mM
EDTA) and sonicated on ice to disrupt *E. coli*. This was
15 centrifuged at 14,000 r.p.m. at 4°C for 20 minutes to
obtain a precipitate. The precipitate was washed twice
with a cell disruption buffer containing 0.5% Triton X-
100™ and washed with water to remove Triton X-100™. Then,
the resultant mixture was dissolved by soaking in a
20 denaturation buffer containing 8 M urea (8M urea, 50 mM
Tris pH8.5, 20 mM 2ME) at 37°C for 1 hour. The solution
was passed through TALON metal affinity resin (Clontech),
washed with the denaturation buffer containing 10 mM
imidazole, and then eluted with the denaturation buffer
25 containing 100 mM imidazole to purify the solution. The

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purified product was dialyzed against PBS for 3 days with exchanging the buffer every other night to obtain the protein hBSSP5-His or mBSSP5-His.

Example 4

5 Expression of novel serine protease mature protein encoded by BSSP5 gene by using pFBTrypSigTag/hBSSP5

(1) Construction of pFBTrypSigTag/hBSSP5

The sequences represented by SEQ ID NOS: 5 and 6 were subjected to annealing and digested with NheI and BamHI. The resultant fragment was inserted into NheI-BamHI digested pSecTag2A (Invitrogen) to obtain pSecTrypHis. Twenty units of BamHI was added to 5 µg of pSecTrypHis vector and the vector was cleaved at 37°C over 4 hours. Then, 6 units of mung bean nuclease (TAKARA) was added thereto and reacted at room temperature (25°C) for 30 minutes to blunt the terminal ends. Further, the 3'-terminus side of the cloning site was cleaved with 20 units of XhoI, 1 unit of bacterial alkaline phosphatase (TAKARA) was added thereto and the reaction was carried out at 65°C for 30 minutes.

According to the same manner as that described in JP 9-149790 A or Biochim. Biophys. Acta, 1350, 11, 1997, mRNA was prepared from COLO201 cells and cDNA was synthesized to obtain the plasmid pSPORT/neurosin. cDNA of an active region of neurosin was obtained from

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pSPORT/neurosin by PCR using primers having the sequences represented by SEQ ID NOS: 7 and 8. Ten units of XhoI was reacted with the PCR product at 37°C for 3 hours to cleave XhoI site at the 3'-side thereof. This was inserted into
5 pSecTrypHis by TAKARA ligation kit to obtain pSecTrypHis/neurosin (Fig. 3).

Amplification was carried out by using the primers having the sequences represented by SEQ ID NOS: 9 and 10 so that the peptide of Leu-Val-His-Gly was present
10 at the C-terminus of the part from trypsin signal to the enterokinase recognition site of pSecTrypHis/neurosin. This was inserted between NheI and HindIII sites of pSecTag2A to construct the plasmid pTrypSig.

One µg (0.1 µl) of the plasmid pSecTab2A was
15 treated with the restriction enzymes NheI and BamHI to completely remove a region encoding the leader sequence of IgGk. One hundred pmol portions of DANs represented by SEQ ID NOS: 31 and 32 were added to the resultant solution and the mixture was heated at 70°C for 10 minutes and subjected
20 to annealing by allowing to stand at room temperature for 30 minutes. Two µl of I solution of DNA ligation kit Ver. 2 (TAKARA) was added to 1 µl portions of His secretory signal sequence and pSecTag2A treated by NheI and BamHI and the reaction was carried out at 16°C for 30 minutes.

25 To the reaction mixture was add 0.1 ml of *E. coli*

competent cell XL1-Blue (STRATAGENE) and reacted on ice for 30 minutes. Then, the reaction mixture was subjected to heat shock at 42°C for 60 seconds. After standing on ice for 2 minutes, 0.9 ml of SOC culture medium (Toyo Boseki K.K.) was added thereto and the mixture was shaken with a shaker at 37°C for 1 hour. The mixture was centrifuged at 5,000 r.p.m. for 1 minutes and the supernatant was discarded. The precipitated competent cells were suspended in the liquid remained in the centrifuge tube and the suspension was applied to 2 ampicillin LB plates containing 100 µg/ml of ampicillin in the ratio of 1 : 10. The plates were incubated at 37°C overnight. Among the colonies formed, a colony into which DNA of His secretory signal was inserted was selected by PCR to obtain pTrypHis.

A sequence of about 200 bp containing His Tag region of pTrypHis was amplified by using primers having the sequence represented by SEQ ID NOS: 10 and 11 and a fragment of about 40 bp containing His Tag and enterokinase recognizing site formed by digestion of HindIII and BamHI was inserted into pTrypSig to construct pTrypSigTag (Fig. 4A).

cDNA was prepared by PCR of the sequence from trypsin signal to enterokinase recognizing site of pTrypSigTag using primers having the sequences represented by SEQ ID NOS 8 and 12 and cut out by digestion with BglII

and BamHI. It was inserted into BamHI site of pFastBAC1 (GIBCO). The insertion direction was confirmed by PCR using primers having the sequences represented by SEQ ID NOS 8 and 13. A clone into which the cDNA was inserted in the direction toward transcription and translation by polyhedrin promoter was selected to obtain pFBTrypSigTag.

Twenty units of BamHI was added to 5 µg of pFBTrypSigTag vector and the vector was cleaved at 37°C over 4 hours, followed by addition of 6 units of mung bean nuclease (TAKARA) and reaction at room temperature (25°C) for 30 minutes to blunt the terminal ends. Further, the 3'-side of the cloning site was cleaved by 20 units of EcoRI, followed by addition of 1 unit of bacterial alkaline phosphatase (TAKARA). The reaction was carried out at 65°C for 30 minutes.

cDNA of the active region of hBSSP5 was obtained from pTrcHis/hBSSP5 prepared from E. coli pTrcHis/hBSSP5 (accession No. FERM P-17038) or pCRII/hBSSP5 by PCR according to a conventional manner. The resultant cDNA was inserted into pFBTrypSigTag to obtain pFBTrypSigTag/hBSSP5 (Fig. 4B). At this time, correct insertion of hBSSP5 was confirmed by determining the sequence.

Bacmid DNA was transformed with pFBTrypSigTag/hBSSP5 according to a protocol of Gibco BRL BAC-TO-BAC baculovirus expression system to prepare a

recombinant bacmid having chimera hBSSP5 fused with trypsinogen signal peptide, His tag and enterokinase recognizing site. When this was expressed in Sf-9 cell according to a manual of BAC-TO-BAC baculovirus expression system, it was secreted in the culture supernatant from 2 days after infection of the virus.

According to the same manner as described above, pFBTrypSigTag/mBSSP5 can be prepared and secreted by using pTrcHis/mBSSP5 obtained from E. coli pTrcHis/mBSSP5 (accession No. FERM P-17035) or pCRII/mBSSP5 obtained in Example 1.

(2) Determination of enzyme activity

The recombinant fused protein hBSSP5 obtained in the culture supernatant was passed through a chelate column to purify it and, after dialysis, its enzyme activity was determined. First, the culture supernatant was applied to a chelate column (Ni-NTA-Agarose, Qiagen) with PBS buffer and eluted stepwise with a solution of imidazole (Wako Pure Chemical Industries, Ltd.) dissolved in PBS. The resultant imidazole-eluted fraction was applied to a PD-10 column (Pharmacia) to exchange to PBS buffer. Fifty μ l of this sample was mixed with 10 μ l of enterokinase (1 U/1 μ l, Invitrogen) and the reaction was carried out at room temperature for 60 minutes. Each of various synthetic substrates (Peptide Laboratory, Boc-Gln-Ala-Arg-MCA, Boc-

Phe-Ser-Arg-MCA, Bz-Arg-MCA, Boc-Val-Leu-Lys-MCA, Pyr-Gly-Arg-MCA, Pro-Phe-Arg-MCA, Boc-Val-Pro-Arg-MCA, Z-Arg-Arg-MCA, Arg-MCA, Z-Phe-Arg-MCA) was dissolved in DMSO and diluted with 1 M Tris-HCl (pH 8.0) to obtain a substrate solution. Fifty μ l of 0.2 M substrate solution was added thereto and further the reaction was carried out at 37°C. After one hour, the fluorescence of AMC (7-amino-4-methylcoumalin) formed by the enzymatic reaction was measured at 380 nm of excitation wavelength and 460 nm of fluorescence wavelength to determine the activity.

As a result, the recombinant fused protein hBSSP5 has been shown to have serine protease activity. Likewise, mBSSP5 derived from a mouse showed the activity.

Example 5

Detection of BSSP5 in urine

Urine samples were collected from human beings and rats. According to a conventional method, 10 μ l of each sample was subjected to electrophoresis on 12.5% SDS-polyacrylamide gel and then blotted on PVDF membrane (1 mmobilon P, Millipore). After blocking the filter with skimmed milk, and reacted with an anti-BSSP5 antibody diluted 1,000-fold or 10,000-fold with Tween-PBS at room temperature for several hours to overnight. The filter was washed with Tween-PBS three times and reacted with alkaline phosphatase labeled anti-rabbit IgG, followed by washing

Tween-PBS in the same manner. When the filter was colored by dipping in BCIP/NBT solution, BSSP5 bands having the presumed molecular weight were detected in the human benign and rat samples (Fig. 5).

5 The anti-BSSP5 antibody was a peptide antibody against hBSSP5. The antibody was prepared as follows.

 Namely, peptides were synthesized by adding one cysteine to C-terminus sides of the peptides composed of the 56th to 73rd amino acids (Glu Tyr Asp Arg Ser Ser Asn Ala Glu Pro Leu Gln Val Leu Ser Val Ser Arg) and the 207th to 225th amino acids (Asn Val Arg Ala Pro Ala Val Tyr Thr Arg Val Ser Lys Phe Ser Thr Trp Ile Asn) of SEQ ID NO: 2. Separately, hemocyanin (KLH) was reacted with a cross-linking agent, m-maleimidobenzoyl-N-hydroxysuccinimide ester (MBS) to prepare a KLH-MB complex. The KLH-MB was reacted each synthetic peptide to obtain two immunogens. The resultant immunogen was administered to a rabbit together with Freund's complete adjuvant once. Then, booster immunization was conducted together with Freund's incomplete adjuvant every two weeks three times in all. Four days after the last booster immunization, a blood sample was collected. The serum obtained was purified by protein A column to obtain a peptide antibody against hBSSP5.

25 Reactivity of each peptide antibody to BSSP5 was

confirmed. The anti-BSSP5 antibody used in Example 5 and the following Example 6 is a mixture of these two peptide antibodies.

Example 6

5 Variation in blood BSSP5 in rat pancreatitis model

 According to a conventional manner, pancreatitis was induced in 4 to 6-week-old rats with cerulein. Blood samples were collected before inducing pancreatitis and 6,
10 12 and 24 hours after inducing pancreatitis to recover serum samples. Serum albumin was removed by mixing with Blue Sepharose (Amersham Pharmacea) and, according to the same manner as in Example 1, 10 μ l of the serum was subjected to SDS-PAGE western blotting, followed by
15 detection with the anti-BSSP 5 antibody. As a result, although BSSP5 was present even in a healthy state, 12 hours after inducing pancreatitis, temporary increase in blood BSSP5 was observed and this showed the possibility of detection of pancreatitis by measuring blood BSSP5.

20

INDUSTRIAL UTILITY

 According to the present invention, there are provided isolated human and mouse serine protease (hBSSP5 and mBSSP5) polynucleotides, their homologous forms, mature
25 forms, precursors and polymorphic variants. Further,

according to the present invention, there are provided hBSSP5 and mBSSP5 proteins as well as compositions containing hBSSP5 and mBSSP5 polynucleotides and proteins, their production and use.

5

SEQUENCE LISTING FREE TEXT

SEQ ID NO: 5: Designed oligonucleotide to construct plasmid pSecTrypHis

10 SEQ ID NO: 6: Designed oligonucleotide to construct plasmid pSecTrypHis

SEQ ID NO: 7: Designed oligonucleotide primer to amplify neurosin-encoding sequence

SEQ ID NO: 8: Designed oligonucleotide primer to amplify neurosin-encoding sequence

15 SEQ ID NO: 9: Designed oligonucleotide primer to amplify a portion of plasmid pSecTrypHis/Neurosin

SEQ ID NO: 10: Designed oligonucleotide primer to amplify a portion of plasmid pSecTrypHis/Neurosin

20 SEQ ID NO: 11: Designed oligonucleotide primer to amplify a portion of plasmid pTrypHis

SEQ ID NO: 12: Designed oligonucleotide primer to amplify a portion of plasmid pTrypSigTag

SEQ ID NO: 13: Designed oligonucleotide primer to amplify a portion of plasmid pFBTrypSigTag

25 SEQ ID NO: 14: Designed oligonucleotide primer to

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amplify conserved region of serin proteases-encoding sequence; n is a, c, g or t.

SEQ ID NO: 15: Designed oligonucleotide primer to amplify conserved region of serin proteases-encoding
5 sequence; n is a, c, g or t.

SEQ ID NO: 16: Designed oligonucleotide primer for RACE for hBSSP5 (forward)

SEQ ID NO: 17: Designed oligonucleotide primer for RACE for hBSSP5 (forward)

10 SEQ ID NO: 18: Designed oligonucleotide primer for RACE for hBSSP5 (reverse)

SEQ ID NO: 19: Designed oligonucleotide primer for RACE for hBSSP5 (reverse)

15 SEQ ID NO: 20: Designed oligonucleotide primer designated as hBSSP5F1 to amplify full length hBSSP5 (forward)

SEQ ID NO: 21: Designed oligonucleotide primer designated as hBSSP5F2 to amplify mature hBSSP5-encoding region (forward)

20 SEQ ID NO: 22: Designed oligonucleotide primer designated as hBSSP5R1/E to amplify full length hBSSP5 (reverse)

SEQ ID NO: 23: Designed oligonucleotide primer designated as hBSSP5R4 for RT-PCR (reverse)

25 SEQ ID NO: 24: Designed oligonucleotide primer

designated as mBSSP5F1 for RACE for mBSSP5 (forward)

SEQ ID NO: 25: Designed oligonucleotide primer
designated as mBSSP5F2 for RACE for mBSSP5 (Forward)

5 SEQ ID NO: 26: Designed oligonucleotide primer
designated as mBSSP5F3 to amplify full length mBSSP5
(forward)

SEQ ID NO: 27: Designed oligonucleotide primer
designated as mBSSP5Fmature to amplify mature mBSSP5-
encoding region (forward)

10 SEQ ID NO: 28: Designed oligonucleotide primer
designated as mBSSP5.1 for RACE for mBSSP5 (reverse)

SEQ ID NO: 29: Designed oligonucleotide primer
designated as mBSSP5R2 for RACE for mBSSP5 (reverse)

15 SEQ ID NO: 30: Designed oligonucleotide primer
designated as mBSSP5R3/E to amplify full length mBSSP5
(reverse)

SEQ ID NO: 31: Designed oligonucleotide to
construct plasmid pTrypHis

20 SEQ ID NO: 32: Designed oligonucleotide to
construct plasmid pTrypHis

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What is claimed is:

1. A protein having the amino acid sequence composed of 231 amino acids represented by the 1st to 231st amino acids of SEQ ID NO: 2; or a protein having an amino acid sequence derived from the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2 by deletion, substitution or addition of one to several amino acids and having the same property as that of the protein having the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2; or a modified derivative thereof.

2. A nucleotide sequence represented by the 110th to 802nd bases of SEQ ID NO: 1; a nucleotide sequence encoding the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2; or a nucleotide sequence hybridizable with a nucleotide sequence which is complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein having the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 2.

3. A protein having the amino acid sequence composed of 231 amino acids represented by the 1st to 231st amino acids of SEQ ID NO: 4; or a protein having an amino acid sequence derived from the amino acid sequence

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represented by the 1st to 231st amino acids of SEQ ID NO: 4 by deletion, substitution or addition of one to several amino acids and having the same property as that of the protein having the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 4; or a modified derivative thereof.

4. A nucleotide sequence represented by the 132nd to 824th bases of SEQ ID NO: 3; a nucleotide sequence encoding the amino acid sequence represented by SEQ ID NO: 4; or a nucleotide sequence hybridizable with a nucleotide sequence which is complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein having the amino acid sequence represented by the 1st to 231st amino acids of SEQ ID NO: 4.

5. A protein having the amino acid sequence composed of 33 amino acids represented by the -33rd to -1st amino acids of SEQ ID NO: 4; or a protein having an amino acid sequence derived from the amino acid sequence represented by the -33rd to -1st amino acids of SEQ ID NO: 4 by deletion, substitution or addition of one to several amino acids and having the same property as that of the protein having the amino acid sequence represented by the -33rd to -1st amino acids of SEQ ID NO: 4; or a modified derivative or fragment thereof.

6. A nucleotide sequence represented by the 33rd to 131st bases of SEQ ID NO: 3; a nucleotide sequence encoding the amino acid sequence represented by the -33rd to -1st amino acids of SEQ ID NO: 4; or a nucleotide sequence hybridizable with a nucleotide sequence which is complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein having the amino acid sequence represented by the -33rd to -1st amino acids of SEQ ID NO: 4; or a fragment thereof.

7. A protein having the amino acid sequence of 264 amino acids represented by the -33rd to 231st amino acids of SEQ ID NO: 4; or a protein having an amino acid sequence derived from the amino acid sequence represented by the -33rd to 231st amino acids of SEQ ID NO: 4 by deletion, substitution or addition of one to several amino acids and having the same property as that of the protein having the amino acid sequence represented by the -33rd to 231st amino acids of SEQ ID NO: 4; or a modified derivative thereof.

8. A nucleotide sequence represented by the 33rd to 824th bases of SEQ ID NO: 3; a nucleotide sequence encoding the amino acid sequence represented by the -33rd to 231st amino acids of SEQ ID NO: 4; or a nucleotide sequence hybridizable with a nucleotide sequence which is

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complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein having the amino acid sequence represented by the -33rd to 231st amino acids of SEQ ID NO: 4.

9. A nucleotide sequence represented by SEQ ID NO: 1; or a nucleotide sequence hybridizable with a nucleotide sequence which is complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein encoded by the nucleotide sequence represented by SEQ ID NO: 1.

10. A nucleotide sequence represented by SEQ ID NO: 3; or a nucleotide sequence hybridizable with a nucleotide sequence which is complementary to the above nucleotide sequence under stringent conditions and encoding a protein having the same property as that of the protein encoded by the nucleotide sequence represented by SEQ ID NO: 3.

11. A vector comprising the nucleotide sequence according to any one of claims 2, 4, 6 and 8-10.

12. Transformed cells having the nucleotide sequence according to any one of claims 2, 4, 6 and 8-10 in an expressible state.

13. A process for producing a protein which

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comprises culturing cells transformed with the nucleotide sequence according to either of claims 2 and 9, and collecting hBSSP5 produced.

5 14. A process for producing a protein which comprises culturing cells transformed with the nucleotide sequence according to any one of claims 4, 6, 8 and 10, and collecting mBSSP5 produced.

10 15. The process according to claim 13 or 14, wherein the cells are *E. coli* cells, animal cells or insect cells.

16. A non-human transgenic animal whose expression level of BSSP5 gene has been altered.

15 17. The non-human transgenic animal according to claim 16, wherein BSSP5 gene is cDNA, genomic DNA or synthetic DNA encoding BSSP5.

18. The non-human transgenic animal according to claim 16, wherein the expression level has been altered by mutating a gene expression regulatory site.

20 19. A knockout mouse whose mBSSP5 gene function is deficient.

20. An antibody against the protein according to any one of claims 1, 3, 5 and 7 or a fragment thereof.

25 21. The antibody according to claim 20 which is a polyclonal antibody, a monoclonal antibody or a peptide antibody.

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22. A process for producing a monoclonal antibody against the protein according to any one of claims 1, 3, 5 and 7 or a fragment thereof which comprises administering the protein according to any one of claims 1, 3, 5 and 7 or a fragment thereof to a warm-blooded animal other than a human being, selecting the animal whose antibody titer is recognized, collecting its spleen or lymph node, fusing the antibody producing cells contained therein with myeloma cells to prepare a monoclonal antibody producing hybridoma.

23. A method for determining the protein according to any one of claims 1, 3, 5 and 7 or a fragment thereof in a specimen which is based on immunological binding of an antibody against the protein or a fragment thereof to the protein or a fragment thereof.

24. A method for determining hBSSP5 or a fragment thereof in a specimen which comprises reacting a monoclonal antibody or a polyclonal antibody against the protein according to claim 1 or a fragment thereof and a labeled antibody with hBSSP5 or a fragment thereof in the specimen to detect a sandwich complex produced.

25. A method for determining hBSSP5 or a fragment thereof in a specimen which comprises reacting a monoclonal antibody or a polyclonal antibody against the protein according to claim 1 or a fragment thereof with

labeled hBSSP5 and hBSSP5 or a fragment thereof in the specimen competitively to detect an amount of hBSSP5 or a fragment thereof in the specimen based on an amount of the labeled hBSSP5 reacted with the antibody.

5 26. The method according to any one of claims 23-25, wherein the specimen is a body fluid.

 27. A diagnostic marker for diseases in tissues comprising the protein according to any one of claims 1, 3, 5 and 7.

10 28. The marker according to claim 27 to be used for diagnosis of Alzheimer's disease or epilepsy in brain.

 29. The marker according to claim 27 to be used for diagnosis of cancer or inflammation of brain, prostate, placenta, testicle, spleen or pancreas.

15 30. The marker according to claim 27 to be used for diagnosis of sterility in semen or sperms

 31. The marker according to claim 27 to be used for diagnosis of prostatic hypertrophy in prostate.

20 32. A method for detecting pancreatitis which comprises measuring concentration of the protein according to any one of claims 1, 3, 5 and 7 in blood or urine.

 33. A pharmaceutical composition for detecting pancreatitis which comprises the antibody according to claim 20 or 21.

25 34. Use of the protein according to any one of

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claims 1, 3, 5 and 7 for preparing an antibody for detecting pancreatitis.

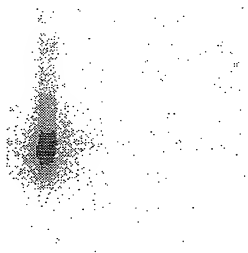
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Fig. 1

hBSSP-5

pancreas
kidney
skeletal muscle
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heart

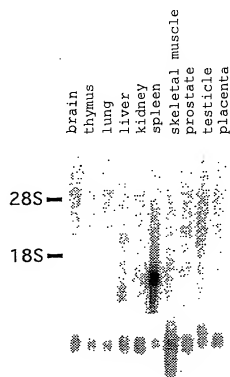


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2/6

Fig. 2

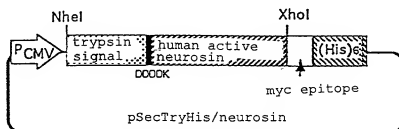
mBSSP-5



09856319.052101

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Fig. 3



09/856319-052101

Fig. 4

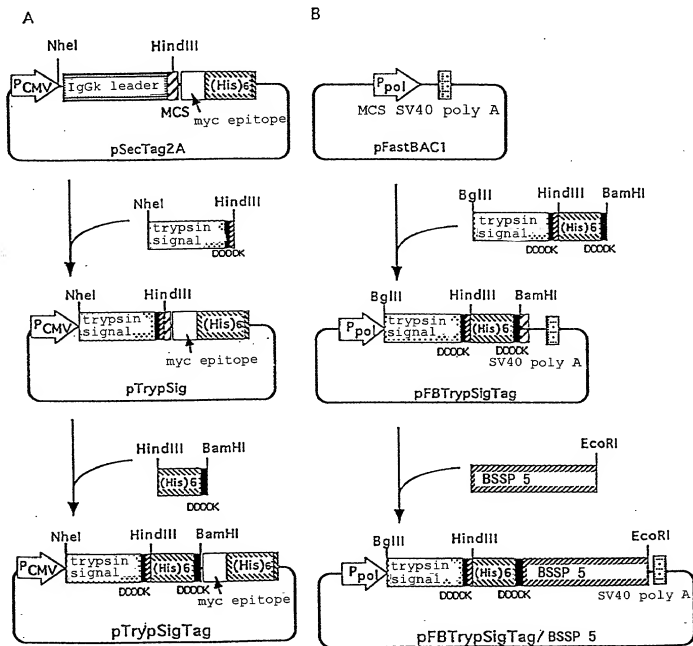


Fig. 5

human being rat



09/856319

6/6

Fig. 6

0, 6, 18, 24 hours



09856319-052101

Combined Declaration for Patent Application and Power of Attorney

As a below-named inventor, I hereby declare that:

My residence, post office address and citizenship are as stated below next to my name; and that I believe I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled

NOVEL SERINE PROTEASE BSSP5

the specification of which (check one)

- [] is attached hereto;
 [] was filed in the United States under 35 U.S.C. §111 on _____, as
 U.S. Appl. No. _____; or
 [x] was/will be filed in the U.S. under 35 U.S.C. §371 by entry into the U.S. national stage of an
 international (PCT) application, PCT/JP9906473; filed 19/11/1999, entry requested on
 _____; national stage application received U.S. Appl. No. _____; §371/§102(e)
 date _____ (* if known)

and was amended on _____ (if applicable).

(include dates of amendments under PCT Art. 19 and 34 if PCT)

I have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment referred to above; and I acknowledge the duty to disclose to the Patent and Trademark Office (PTO) all information known by me to be material to patentability as defined in 37 C.F.R. §1.56.

I hereby claim foreign priority benefits under 35 U.S.C. §§ 119 and 365 of any prior foreign application(s) for patent or inventor's certificate, or prior PCT application(s) designating a country other than the U.S., listed below under the "Yes" box checked and have also identified below any such application having a filing date before that of the application on which priority is claimed.

<u>347806/1998</u>	<u>Japan</u>	<u>20/11/1998</u>	[x]	[]
(Number)	(Country)	(Day Month Year Filed)	YES	NO
_____	_____	_____	[]	[]
(Number)	(Country)	(Day Month Year Filed)	YES	NO

I hereby claim the benefit under 35 U.S.C. §120 of any prior U.S. non-provisional application(s) or prior PCT application(s) designating the U.S. listed below, or under §119(e) of any prior U.S. provisional applications listed below, and, insofar as the subject matter of each of the claims of this application is not disclosed in such U.S. or PCT application in the manner provided by the first paragraph of 35 U.S.C. §112, I acknowledge the duty to disclose to the PTO all information as defined in 37 C.F.R. §1.56(a) which occurred between the filing date of the prior application and the national filing date of this application:

_____ (Application No.)	_____ (Day Month Year Filed)	_____ (Status: patented, pending, abandoned)
_____ (Application No.)	_____ (Day Month Year Filed)	_____ (Status: patented, pending, abandoned)
_____ (Application No.)	_____ (Day Month Year Filed)	_____ (Status: patented, pending, abandoned)

As a named inventor, I hereby appoint the following registered practitioners to prosecute this application and to transact all business in the Patent and Trademark Office connected therewith:

All of the practitioners associated with Customer Number 001444

Direct all correspondence to the address associated with Customer Number 001444; i.e.,

BROWDY AND NEIMARK, P.L.L.C.
 624 Ninth Street, N.W.
 Washington, D.C. 20001-5303
 (202) 628-5197

The undersigned hereby authorizes the U.S. Attorneys or Agents appointed herein to accept and follow instructions from AOYAMA & PARTNERS as to any action to be taken in the U.S. Patent and Trademark Office regarding this application without direct communication between the U.S. Attorneys or Agents and the undersigned. In the event of a change of the persons from whom instructions may be taken, the U.S. Attorneys or Agents appointed herein will be so notified by the undersigned.

00856310, 052701

Title: NOVEL SERINE PROTEASE BSSP5
 U.S. Application filed _____, Serial No. _____
 PCT Application filed Nov. 19, 1999, Serial No. PCT/JP99/06473

I hereby further declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under 18 U.S.C. §1001 and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

1-00	FULL NAME OF FIRST INVENTOR Hidetoshi UEMURA	INVENTOR'S SIGNATURE <i>Hidetoshi Uemura</i>	DATE Apr. 17, 2001
	RESIDENT Itami-shi, Hyogo Japan <i>SPX</i>	CITIZENSHIP Japan	
	POST OFFICE ADDRESS 133, Minamisuzuhara 3-chome, Itami-shi, Hyogo Japan		
2-00	FULL NAME OF SECOND JOINT INVENTOR Akira OKUI	INVENTOR'S SIGNATURE <i>Akira Okui</i>	DATE Apr. 17, 2001
	RESIDENT Yamatokoriyama-shi, Nara Japan <i>SPX</i>	CITIZENSHIP Japan	
	POST OFFICE ADDRESS 295-3, 1000, 1000, 1000, 1000, 1000 Tsutsui-cho, Yamatokoriyama-shi, Nara Japan		
3-00	FULL NAME OF THIRD JOINT INVENTOR Katsuya KOMINAMI	INVENTOR'S SIGNATURE <i>Katsuya Kominami</i>	DATE Apr. 17, 2001
	RESIDENT Hannan-shi, Osaka Japan <i>SPX</i>	CITIZENSHIP Japan	
	POST OFFICE ADDRESS 786-2, Jinenda, Hannan-shi, Osaka Japan		
4-00	FULL NAME OF FOURTH JOINT INVENTOR Nozomi YAMAGUCHI	INVENTOR'S SIGNATURE <i>Nozomi Yamaguchi</i>	DATE Apr. 17, 2001
	RESIDENT Kyoto-shi, Kyoto Japan <i>SPX</i>	CITIZENSHIP Japan	
	POST OFFICE ADDRESS 285-79, Shingoryoguchi-cho, Teramachinishi-iru, Kuramaguchi-dori, Kita-ku, Kyoto-shi, Kyoto Japan		
5-00	FULL NAME OF FIFTH JOINT INVENTOR Shinichi MITSUI	INVENTOR'S SIGNATURE <i>Shinichi Mitsui</i>	DATE Apr. 17, 2001
	RESIDENT Kyoto-shi, Kyoto Japan <i>SPX</i>	CITIZENSHIP Japan	
	POST OFFICE ADDRESS 202, Kitashirakawa-koporasu, 86, Kitashirakawanishi-machi, Sakyo-ku, Kyoto-shi, Kyoto Japan		
	FULL NAME OF SIXTH JOINT INVENTOR	INVENTOR'S SIGNATURE	DATE
	RESIDENT	CITIZENSHIP	
	POST OFFICE ADDRESS		
	FULL NAME OF SEVENTH JOINT INVENTOR	INVENTOR'S SIGNATURE	DATE
	RESIDENT	CITIZENSHIP	
	POST OFFICE ADDRESS		

ALL INVENTORS MUST REVIEW APPLICATION AND DECLARATION BEFORE SIGNING. ALL ALTERATIONS MUST BE INITIALED AND DATED BY ALL INVENTORS PRIOR TO EXECUTION. NO ALTERATIONS CAN BE MADE AFTER THE DECLARATION IS SIGNED. ALL PAGES OF DECLARATION MUST BE SEEN BY ALL INVENTORS.

SEQUENCE LISTING

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00556310-05101

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<211> 27

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<223> Designed oligonucleotide to construct plasmid pTrypHis

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<223> Designed oligonucleotide to construct plasmid pTrypHis

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